

# Principle Components Analysis (PCA) and Face Recognition

Computer Vision (CS0029)

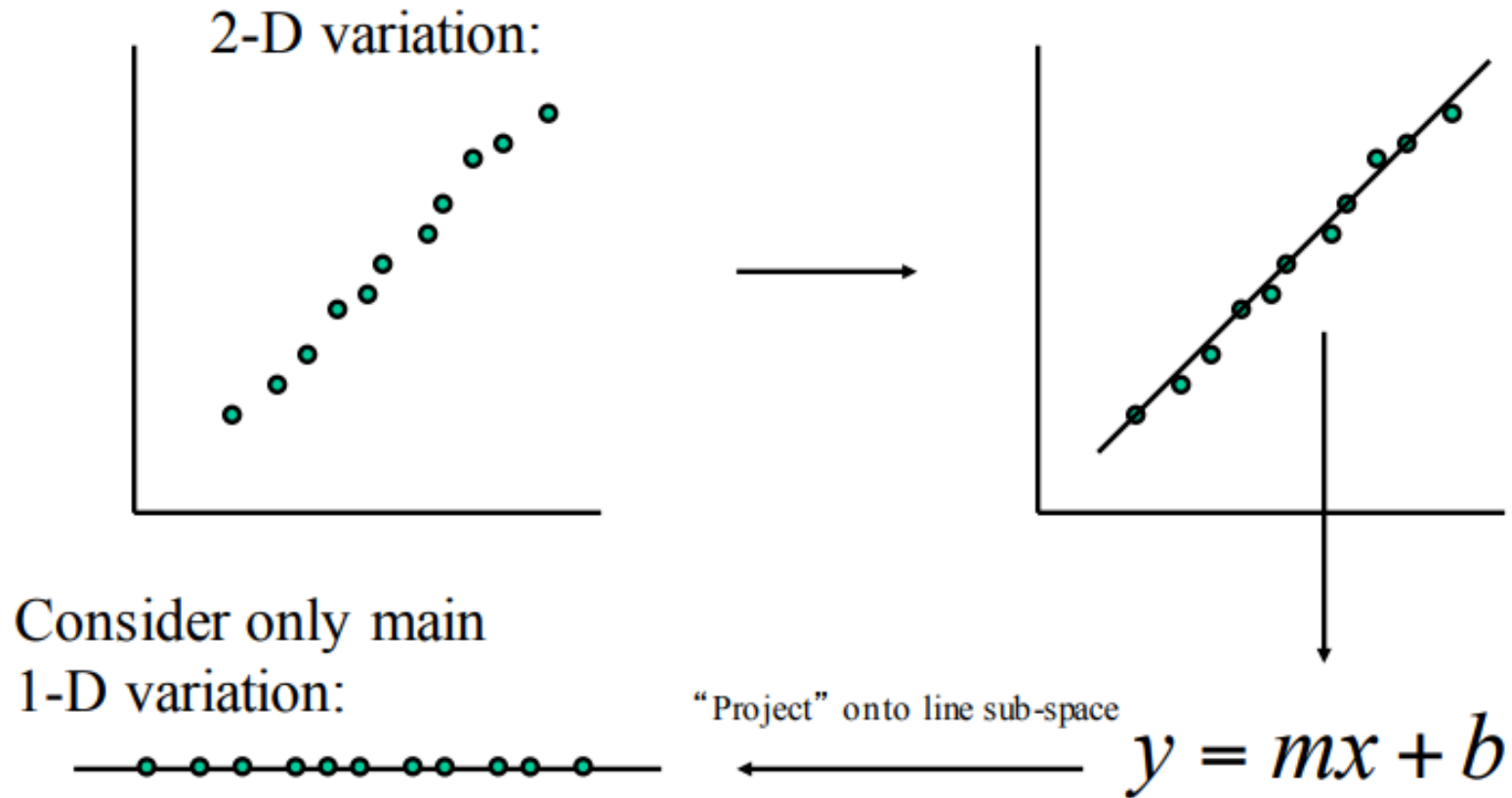
# Feature Sub-Space

- A high-dimensional feature vector is often contained within a lower-dimensional “sub-space”
- When processing the feature data (for modeling or recognition), it is beneficial to deal with the lower-dimensional sub-space
  - Modeling speed, noise
- PCA offers linear approximation to the sub-space which can be reduced to only the major sub-space dimensions
  - The dimensions of the sub-space has high data variance

# Outline (PCA)

- **The most basic linear algebra**
  - **Linear Basis Set**
- Connect PCA to Gaussian Distribution (Ellipse shape distribution in 2D)
- Eigenvalue and eigenvector
- Connect eigenvalue and eigenvector to ellipse
  - Ellipse contour in 2D
- Extend ellipse to Gaussian
  - Covariance matrix
  - Gaussian contours
- PCA face recognition

# Dimensionality Reduction

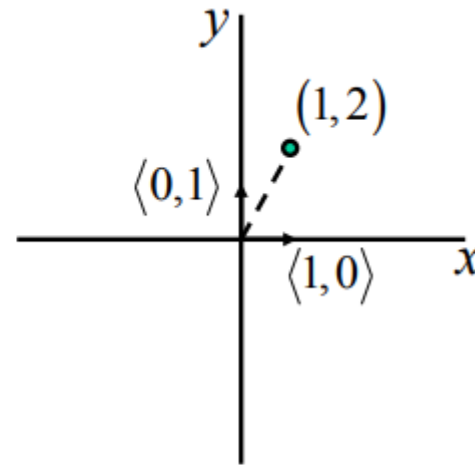


# Linear Basis Set

- 2-D basis set
  - Coordinate is weights of bases

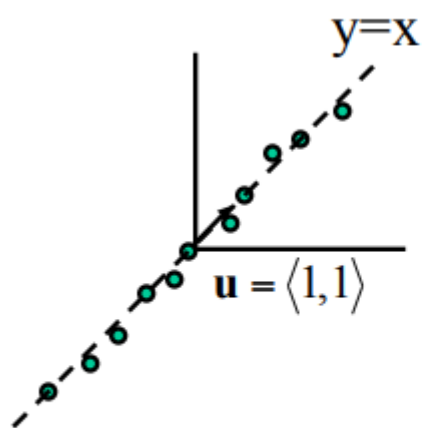
$$\mathbf{x} = \begin{bmatrix} x \\ y \end{bmatrix} = \begin{bmatrix} 1 \\ 2 \end{bmatrix} = 1 \cdot \begin{bmatrix} 1 \\ 0 \end{bmatrix} + 2 \cdot \begin{bmatrix} 0 \\ 1 \end{bmatrix}$$

$\begin{array}{ccc} \uparrow & \nearrow & \uparrow \\ \text{x,y} & & \text{x-axis} \quad \text{y-axis} \\ \text{coordinates} & & \end{array}$



# Linear Basis Set

- 1-D sub-space basis set



2-D space:

$$\mathbf{x}_i = a_i \cdot \begin{bmatrix} 1 \\ 0 \end{bmatrix} + b_i \cdot \begin{bmatrix} 0 \\ 1 \end{bmatrix}$$

1-D sub-space:

$$y_i = \gamma_i \cdot \mathbf{u} = \gamma_i \cdot \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

# Linear Basis Set

- Generate smaller dimension basis for feature vectors

- $x_i = \gamma_1 u_1 + \gamma_2 u_2 + \gamma_3 u_3 + \dots + \gamma_m u_m$

where  $\dim(x) > m$

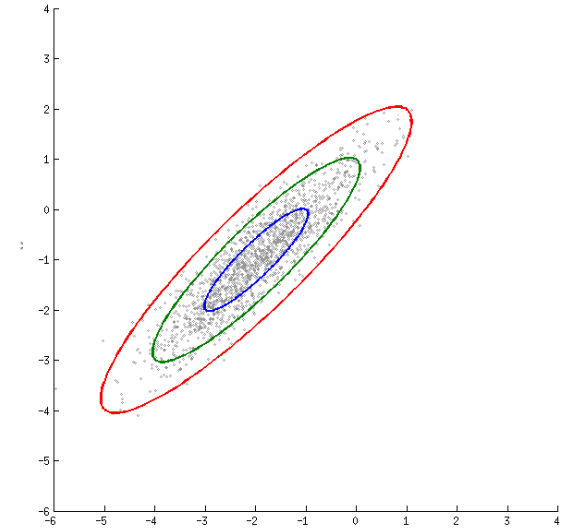
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# Principal Components Analysis (PCA)

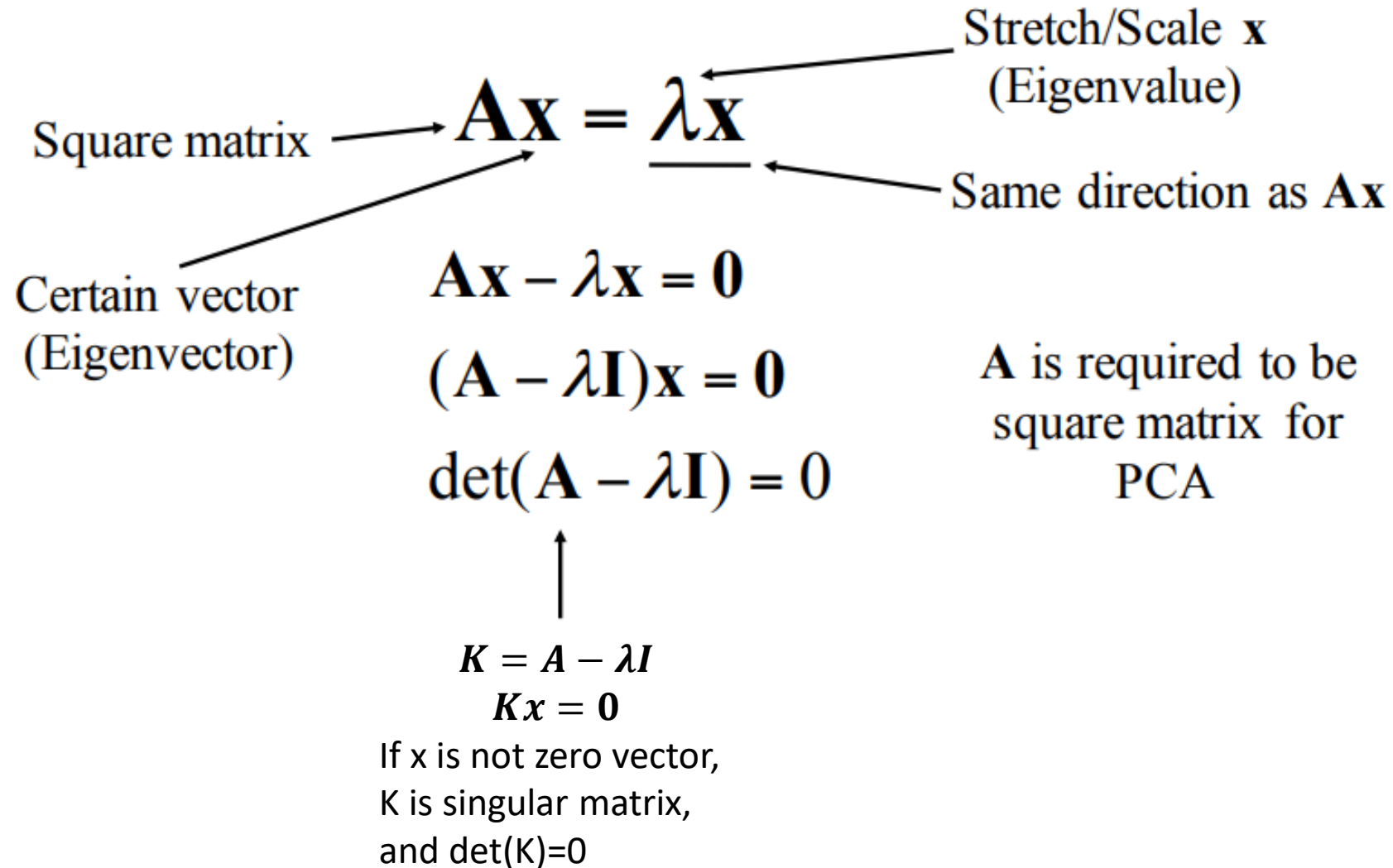
- The main idea for PCA
  - Fit a multi-dimensional Gaussian around data
    - Assumption: data distribution is close to a Gaussian distribution
    - Use covariance of data to model Gaussian
- Select only those dimensions capturing most of the variance in data
  - Reduce dimensionality



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# PCA Primer: Equation for Eigenvalues



# PCA Primer: Equation for Eigenvalues

$$\mathbf{A} = \begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix}$$

$$\mathbf{A} - \lambda \mathbf{I} = \begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} - \begin{bmatrix} \lambda & 0 \\ 0 & \lambda \end{bmatrix} = \begin{bmatrix} 1-\lambda & 2 \\ 2 & 4-\lambda \end{bmatrix}$$

$$\det(\mathbf{A} - \lambda \mathbf{I}) = (1-\lambda)(4-\lambda) - (2)(2) = \lambda^2 - 5\lambda = 0$$

two roots:  $\lambda_1 = 0$ ,  $\lambda_2 = 5$

# PCA Primer: Equation for Eigenvalues

$$(\mathbf{A} - \lambda_i \mathbf{I}) \mathbf{x}_i = \mathbf{0}$$

$$(\mathbf{A} - \lambda_1 \mathbf{I}) \mathbf{e}_1 = \mathbf{0}$$

$$\left( \begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} - \begin{bmatrix} \textcircled{0} & 0 \\ 0 & \textcircled{0} \end{bmatrix} \right) \mathbf{e}_1 = \mathbf{0}$$

$$\begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} \begin{bmatrix} x_1 \\ y_1 \end{bmatrix} = \mathbf{0}$$

$$\mathbf{e}_1 = \begin{bmatrix} x_1 \\ y_1 \end{bmatrix} = \begin{bmatrix} 2 \\ -1 \end{bmatrix}$$

$$(\mathbf{A} - \lambda_2 \mathbf{I}) \mathbf{e}_2 = \mathbf{0}$$

$$\left( \begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} - \begin{bmatrix} \textcircled{5} & 0 \\ 0 & \textcircled{5} \end{bmatrix} \right) \mathbf{e}_2 = \mathbf{0}$$

$$\begin{bmatrix} -4 & 2 \\ 2 & -1 \end{bmatrix} \begin{bmatrix} x_2 \\ y_2 \end{bmatrix} = \mathbf{0}$$

$$\mathbf{e}_2 = \begin{bmatrix} x_2 \\ y_2 \end{bmatrix} = \begin{bmatrix} 1 \\ 2 \end{bmatrix}$$

# PCA Primer: Test Our Results

$$\mathbf{A}\mathbf{e}_1 = \lambda_1 \mathbf{e}_1$$

$$\begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} \begin{bmatrix} 2 \\ -1 \end{bmatrix} = \begin{bmatrix} (1 \cdot 2 - 2 \cdot 1) \\ (2 \cdot 2 - 4 \cdot 1) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \end{bmatrix} = 0 \begin{bmatrix} 2 \\ -1 \end{bmatrix}$$

$$\mathbf{A}\mathbf{e}_2 = \lambda_2 \mathbf{e}_2$$

$$\begin{bmatrix} 1 & 2 \\ 2 & 4 \end{bmatrix} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = \begin{bmatrix} (1 \cdot 1 + 2 \cdot 2) \\ (2 \cdot 1 + 4 \cdot 2) \end{bmatrix} = \begin{bmatrix} 5 \\ 10 \end{bmatrix} = 5 \begin{bmatrix} 1 \\ 2 \end{bmatrix}$$

# Practice

- $A = \begin{bmatrix} 1 & 2 \\ 3 & 2 \end{bmatrix}$ , calculate eigen values of A?
  - $Ax = \lambda x \Rightarrow Ax - \lambda x = 0 \Rightarrow (A - \lambda I)x = 0$
  - $(A - \lambda I)$  is also a 2x2 matrix and  $\det(A - \lambda I) = 0$  if eigenvector  $x$  is not 0 vector
- Eigenvector ( $x_1$ , and  $x_2$ )
  - Solve  $Ax_1 = \lambda_1 x_1$  and  $Ax_2 = \lambda_2 x_2$

# PCA Primer: Symmetric Matrices

- If  $A$  is symmetric ( $A = A^T$ )
  - Real-valued eigenvalues
  - Eigenvectors can be chosen orthonormal
  - Can be factorized as
    - $A = Q\Lambda Q^T$   
where  $Q$  orthonormal ( $Q^T Q = I$ ) and  $\Lambda$  is diagonal
  - Eigenvalues go into diagonal entries of  $\Lambda$ 
    - Convention: put largest eigenvalues first (descending values)
  - Corresponding orthogonal eigenvectors are normalized (to become orthonormal) and go into columns of  $Q$



# Eigen Decomposition: Numpy

```
import numpy as np

A = np.array([[5, 1], [3, 3]])
eValue, eVector = np.linalg.eig(A) #decompose matrix A
print( "A=\n", A )
print( "\nEigenValues=\n", eValue )
print( "\nEigenVectors(Columns)=\n", eVector )

eigenIndex = 0 #the first eigen value and vector
print("\nEigenIndex: ", eigenIndex)
v = eVector[:,eigenIndex].reshape(2,1) #eigen vector
print("\nv=\n", v)
Av = A.dot(v) #A*v
vx = v*eValue[eigenIndex] #v*x
print("\nAv=\n", Av) #compare A*v = v*x
print("\nvx=\n", vx)

eigenIndex = 1
print("\n\nEigenIndex: ", eigenIndex)
v = eVector[:,eigenIndex].reshape(2,1)
print("\nv=\n", v)
Av = A.dot(v)
vx = v*eValue[eigenIndex]
print("\nAv=\n", Av)
print("\nvx=\n", vx)
```

```
A=
[[5 1]
 [3 3]]

EigenValues=
[6. 2.]

EigenVectors(Columns)=
[[ 0.70710678 -0.31622777]
 [ 0.70710678  0.9486833 ]]

EigenIndex:  0

v=
[[0.70710678]
 [0.70710678]]

Av=
[[4.24264069]
 [4.24264069]]

vx=
[[4.24264069]
 [4.24264069]]

EigenIndex:  1

v=
[[-0.31622777]
 [ 0.9486833 ]]

Av=
[[-0.63245553]
 [ 1.8973666 ]]

vx=
[[-0.63245553]
 [ 1.8973666 ]]
```

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# Positive Definite Matrices

- Matrix  $A$  is positive definite for all non-zero  $x$  if
  - Symmetric and  $x^T A x > 0$

Positive <u>semi</u> -definite if $\mathbf{x}^T \mathbf{A} \mathbf{x} \geq 0$
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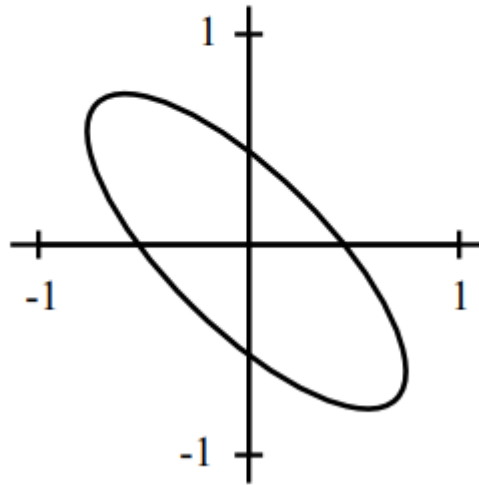
$$\begin{bmatrix} x & y \end{bmatrix} \begin{bmatrix} a & b \\ b & c \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix} > 0$$
$$ax^2 + 2bxy + cy^2 > 0$$

- Recall equation for ellipse
  - Ellipse:  $ax^2 + 2bxy + cy^2 = 1$
  - Center at (0,0)

# Ellipse Factorization

- Consider the rotated ellipse:
- Know the shape of ellipse?
  - Calculate axes and length

$$5x^2 + 8xy + 5y^2 = 1$$

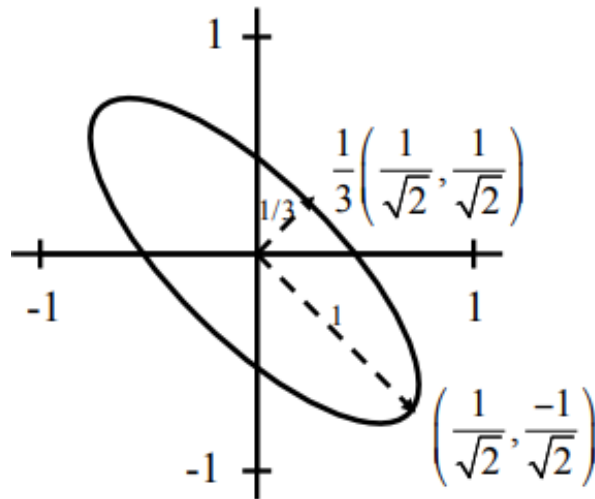


# Ellipse Factorization

$$5x^2 + 8xy + 5y^2 = 1 \rightarrow a = 5, b = 4, c = 5$$

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = 1$$

$$\mathbf{A} = \begin{bmatrix} 5 & 4 \\ 4 & 5 \end{bmatrix} = \mathbf{Q} \mathbf{\Lambda} \mathbf{Q}^T = \begin{bmatrix} 1 & 1 \\ 1 & -1 \end{bmatrix}_{\sqrt{2}} \begin{bmatrix} 9 & 0 \\ 0 & 1 \end{bmatrix} \begin{bmatrix} 1 & 1 \\ 1 & -1 \end{bmatrix}_{\sqrt{2}}$$



Axes of ellipse point along eigenvectors

Half-lengths of axes are  $1/\sqrt{\lambda_i}$   
(NOTE: bigger eigenvalues give shorter axes!)

# Eigenvector projection

- The matrix  $Q^T$  acts as a rotation matrix

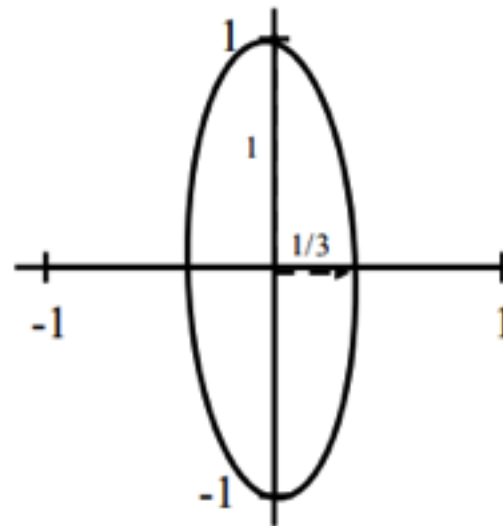
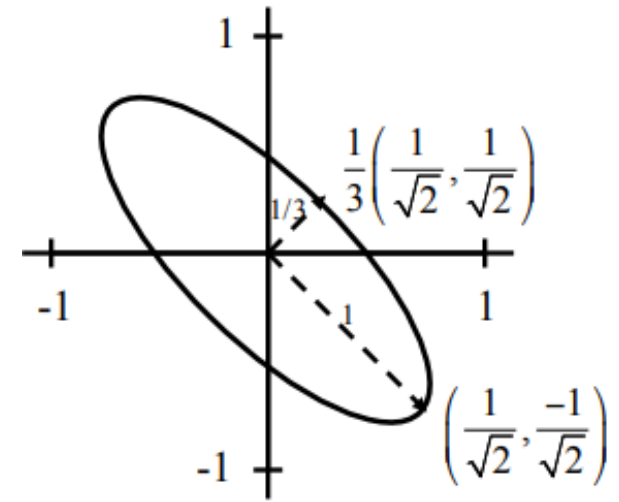
$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \mathbf{x}^T (\mathbf{Q} \mathbf{\Lambda} \mathbf{Q}^T) \mathbf{x} = (\mathbf{x}^T \mathbf{Q}) \mathbf{\Lambda} (\mathbf{Q}^T \mathbf{x}) = \mathbf{X}^T \mathbf{\Lambda} \mathbf{X}$$

$$\mathbf{X} = \mathbf{Q}^T \mathbf{x} = \begin{bmatrix} \hat{\mathbf{e}}_1^T \mathbf{x} \\ \hat{\mathbf{e}}_2^T \mathbf{x} \end{bmatrix}$$

Rotate previous axes:

$$\mathbf{X}_1 = \mathbf{Q}^T \mathbf{x}_1 = \begin{bmatrix} 1 & 1 \\ 1 & -1 \end{bmatrix}_{\sqrt{2}} \begin{bmatrix} 0 \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ -1 \end{bmatrix}_{\sqrt{2}}$$

$$\mathbf{X}_2 = \mathbf{Q}^T \mathbf{x}_2 = \begin{bmatrix} 1 & 1 \\ 1 & -1 \end{bmatrix}_{\sqrt{2}} \begin{bmatrix} 1 \\ 0 \end{bmatrix} = \begin{bmatrix} 1 \\ 1 \end{bmatrix}_{\sqrt{2}}$$



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# Gaussian Density

$$p(\mathbf{x}) = \frac{1}{(2\pi)^{\frac{n}{2}} |\mathbf{K}|^{\frac{1}{2}}} \cdot e^{\left[-\frac{1}{2}(\mathbf{x}-\mathbf{m})^T \mathbf{K}^{-1}(\mathbf{x}-\mathbf{m})\right]}$$

$$f(x|\mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

Mahalanobis distance:  $(x - m)^T K^{-1}(x - m) = C$

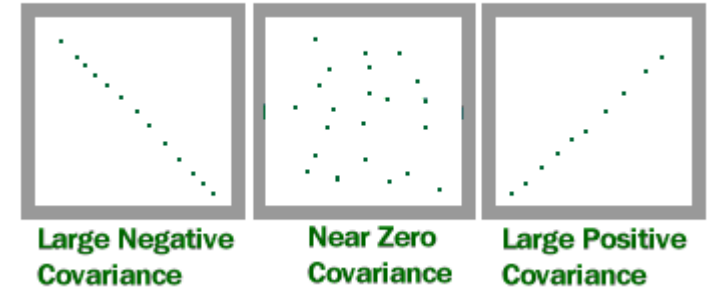
Focus of all points at given distance  $C$  from mean  
(i.e., variance contour with  $C = \#stddev^2$ )

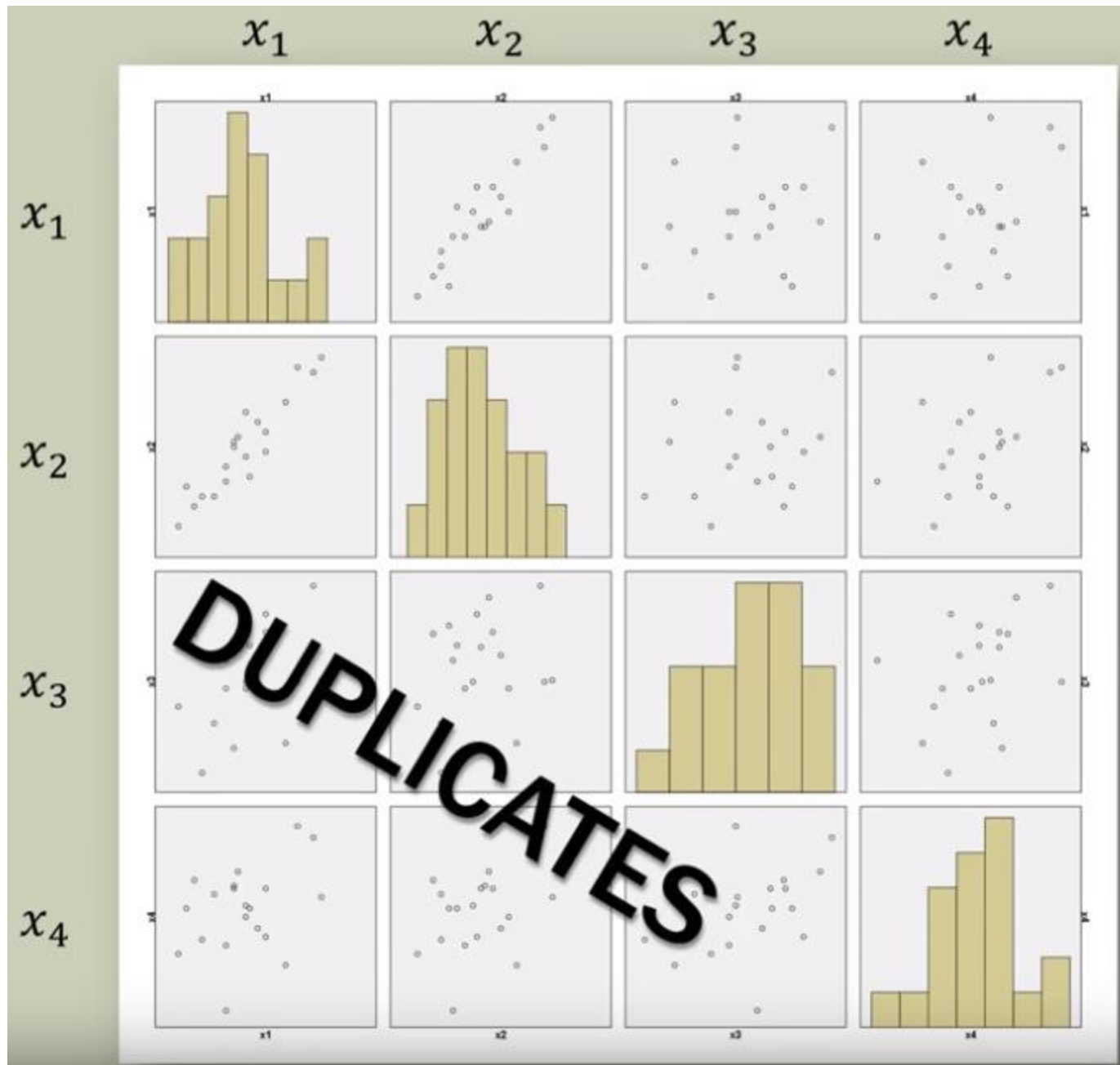


# Covariance

- Covariance is one of the family of statistical measurement used to analyze the linear relationship between two variables.
- Positive covariance: increasing linear relation
- Negative covariance: decreasing linear relation
  - Not strength
- Covariance matrix: covariances from multiple variables
  - symmetric

## COVARIANCE





# Four Variables Covariance Matrix

	$x_1$	$x_2$	$x_3$	$x_4$
$x_1$	$Var(x_1)$	$Cov(x_1, x_2)$	$Cov(x_1, x_3)$	$Cov(x_1, x_4)$
$x_2$		$Var(x_2)$	$Cov(x_2, x_3)$	$Cov(x_2, x_4)$
$x_3$			$Var(x_3)$	$Cov(x_3, x_4)$
$x_4$				$Var(x_4)$

$$\text{Variance: } \text{Var}(X) = \mathbf{E}[(X - \mu)^2]$$

# Covariance Matrix Equation

Covariance matrix ( $K$ ) of  $X$  which is collection of data points:

$K$  is symmetric ( and positive [semi-]definite)

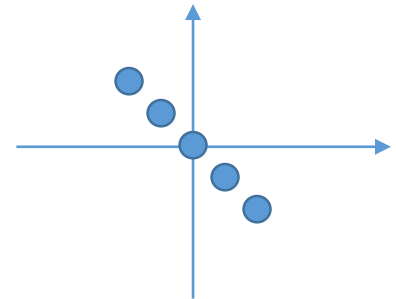
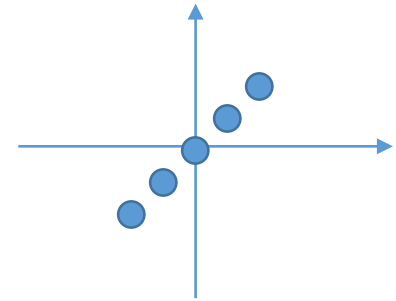
$$\mathbf{X} = (X_1, X_2, \dots, X_n)^T$$

$$K_{X_i X_j} = \text{cov}[X_i, X_j] = \mathbf{E}[(X_i - \mathbf{E}[X_i])(X_j - \mathbf{E}[X_j])]$$

$$K_{\mathbf{X}\mathbf{X}} = \begin{bmatrix} \mathbf{E}[(X_1 - \mathbf{E}[X_1])(X_1 - \mathbf{E}[X_1])] & \mathbf{E}[(X_1 - \mathbf{E}[X_1])(X_2 - \mathbf{E}[X_2])] & \cdots & \mathbf{E}[(X_1 - \mathbf{E}[X_1])(X_n - \mathbf{E}[X_n])] \\ \mathbf{E}[(X_2 - \mathbf{E}[X_2])(X_1 - \mathbf{E}[X_1])] & \mathbf{E}[(X_2 - \mathbf{E}[X_2])(X_2 - \mathbf{E}[X_2])] & \cdots & \mathbf{E}[(X_2 - \mathbf{E}[X_2])(X_n - \mathbf{E}[X_n])] \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{E}[(X_n - \mathbf{E}[X_n])(X_1 - \mathbf{E}[X_1])] & \mathbf{E}[(X_n - \mathbf{E}[X_n])(X_2 - \mathbf{E}[X_2])] & \cdots & \mathbf{E}[(X_n - \mathbf{E}[X_n])(X_n - \mathbf{E}[X_n])] \end{bmatrix}$$

# Example

- Two variables (x, y) case, given 5 data points
  - $(-2,-2), (-1,-1), (0,0), (1,1), (2,2)$
  - $\begin{bmatrix} 5 & 5 \\ 5 & 5 \end{bmatrix}$
- Two variables (x, y) case, given 5 data points
  - $(-2,2), (-1,1), (0,0), (1,-1), (2,-2)$
  - $\begin{bmatrix} 5 & -5 \\ -5 & 5 \end{bmatrix}$



# 3 Variables Covariance Matrix

- 5 data samples

- [-2, 1, 2]
- [-1, 3, 1]
- [0, 5, 0]
- [1, 7, -1]
- [2, 9, -2]

$$\text{Variance: } \text{Var}(X) = \text{E}[(X - \mu)^2]$$

$$\mathbf{X} = (X_1, X_2, \dots, X_n)^T$$

$$K_{X_i X_j} = \text{cov}[X_i, X_j] = \text{E}[(X_i - \text{E}[X_i])(X_j - \text{E}[X_j])]$$

$$K_{\mathbf{X}\mathbf{X}} = \begin{bmatrix} \text{E}[(X_1 - \text{E}[X_1])(X_1 - \text{E}[X_1])] & \text{E}[(X_1 - \text{E}[X_1])(X_2 - \text{E}[X_2])] & \cdots & \text{E}[(X_1 - \text{E}[X_1])(X_n - \text{E}[X_n])] \\ \text{E}[(X_2 - \text{E}[X_2])(X_1 - \text{E}[X_1])] & \text{E}[(X_2 - \text{E}[X_2])(X_2 - \text{E}[X_2])] & \cdots & \text{E}[(X_2 - \text{E}[X_2])(X_n - \text{E}[X_n])] \\ \vdots & \vdots & \ddots & \vdots \\ \text{E}[(X_n - \text{E}[X_n])(X_1 - \text{E}[X_1])] & \text{E}[(X_n - \text{E}[X_n])(X_2 - \text{E}[X_2])] & \cdots & \text{E}[(X_n - \text{E}[X_n])(X_n - \text{E}[X_n])] \end{bmatrix}$$

# 3 Variables Covariance Matrix

- 5 data samples

- [-2, 1, 2]
- [-1, 3, 1]
- [0, 5, 0]
- [1, 7, -1]
- [2, 9, -2]

- Covariance matrix: 
$$\begin{bmatrix} 5 & 4 & -5 \\ 4 & 8 & -4 \\ -5 & 4 & -5 \end{bmatrix}$$

- 5 data samples (3 variable x, y, z)

- $$A = \begin{bmatrix} x & y & z \\ 2 & 1 & 2 \\ -1 & 3 & 1 \\ 0 & 5 & 0 \\ 1 & 7 & -1 \\ 2 & 9 & -2 \end{bmatrix}$$

- $$B = \begin{bmatrix} 2 - \mu_x & 1 - \mu_y & 2 - \mu_z \\ -1 - \mu_x & 3 - \mu_y & 1 - \mu_z \\ 0 - \mu_x & 5 - \mu_y & 0 - \mu_z \\ 1 - \mu_x & 7 - \mu_y & -1 - \mu_z \\ 2 - \mu_x & 9 - \mu_y & -2 - \mu_z \end{bmatrix}$$

- Covariance matrix:  $\frac{(B^T B)}{N}$ 
  - N is number of samples

# Plotting Gaussian Density Function Contours (Physical Meaning of Covariance Matrix)

- $K$  is covariance matrix
  - $K = \frac{(B^T B)}{N}$  (check the previous page)
- Describes a (rotated) ellipse (at a  $C$  variance contour, or squared stdev contour) centered around mean:

$$(x - m)^T K^{-1} (x - m) = C$$

- Thus we can factorize  $K^{-1}$  into eigenvectors (axes) and eigenvalues to give direction and half-lengths of ellipse axes

$$\begin{aligned} K &= Q \Lambda Q^T \\ K^{-1} &= Q \Lambda^{-1} Q^T \end{aligned}$$



# Axes

- To compute the ellipse at Gaussian “variance” contour  $C$  ( $= \#stddev^2$ ):
  - From matrix  $K^{-1}$  (inverse covariance)
    - Axes are columns in  $Q$ ,
    - Half-lengths of axes are  $\frac{\sqrt{C}}{\sqrt{\lambda_i}}$ , with  $\lambda_i$  from  $\Lambda^{-1}$
  - From matrix  $K$  (covariance matrix)
    - Axes are columns in  $Q$
    - Half-lengths of axes are  $\sqrt{C\lambda_i}$ , with  $\lambda_i$  from  $\Lambda$

# Putting it all together

Ellipse  
formula:

$$x^T A x = 1$$

Ellipse defined by  
eigenvectors/values of  $A$

Gaussian &  
Mahalanobis:

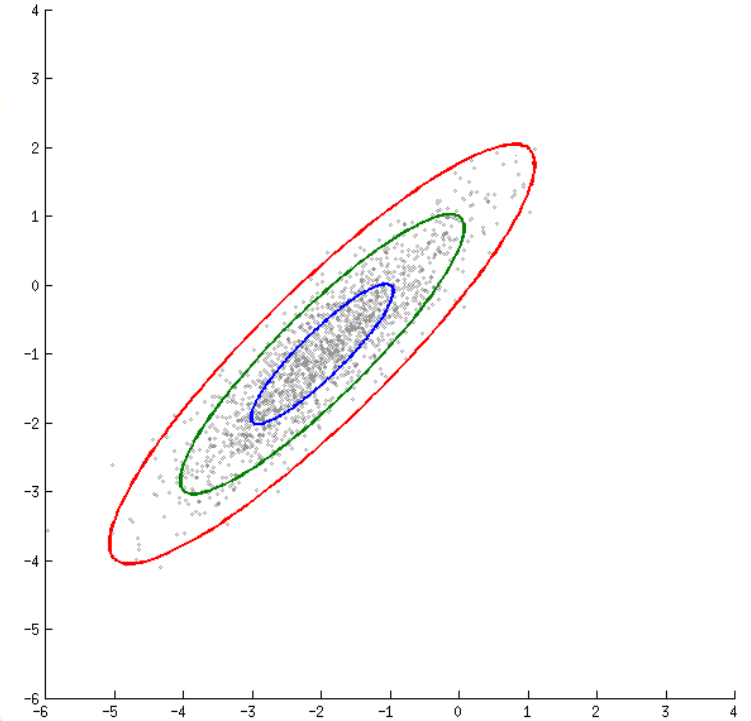
$$(x - m)^T K^{-1} (x - m) = C$$

$$A = K^{-1}$$

Ellipse defined by  
eigenvectors/values of  $K^{-1}$

Larger values of  $C$  give bigger  
variance contour ellipses

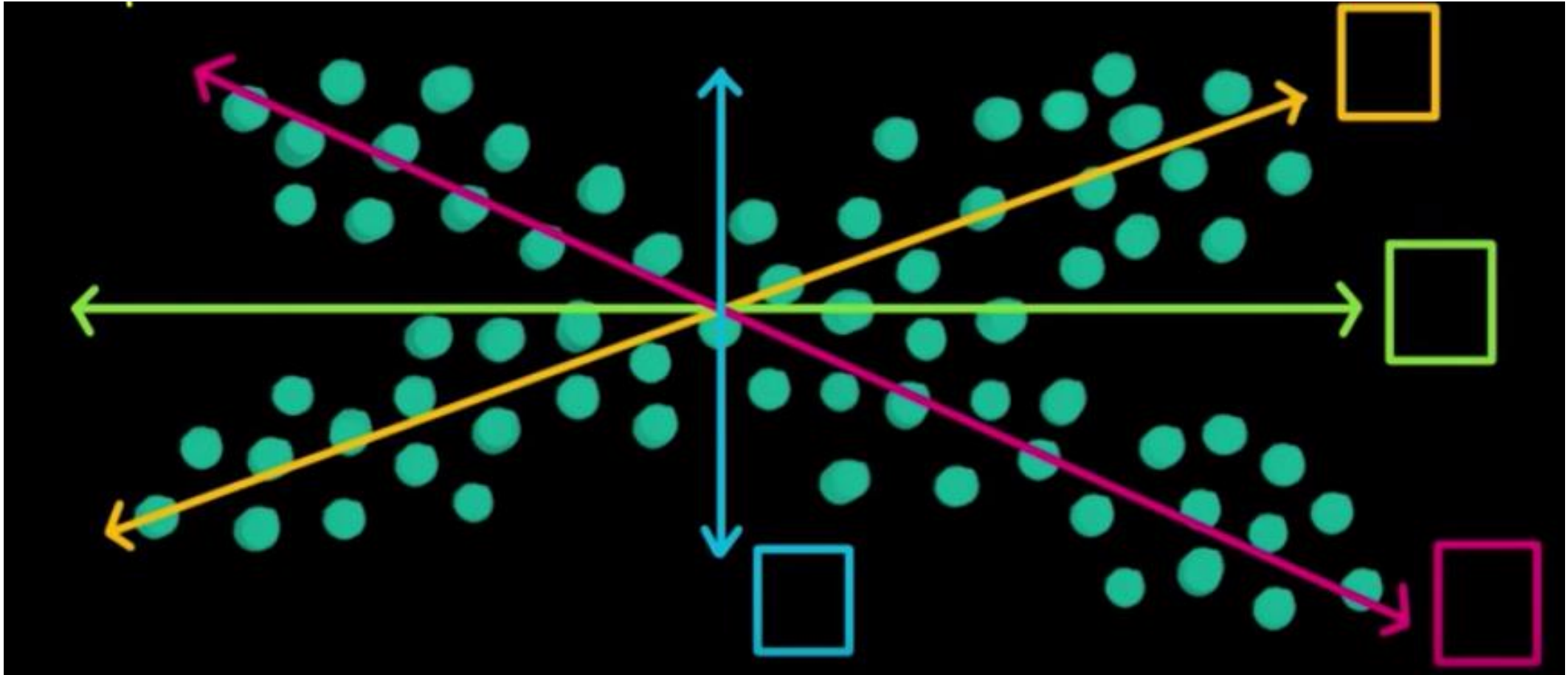
↑  
*Or use  $K$  and just “flip” the  
eigenvalues (saves doing the  
inverse operation)*



# PCA

- Reduce dimensionality
  - Use a Gaussian (covariance matrix) to model a collection of data samples
    - Assumption: data distribution is similar to Gaussian
  - Eigen decomposition to get eigen vector and eigen values
  - Sort eigen vectors by eigen values
  - Use n eigen vectors with largest n eigen values to create a sub-space
  - Project data samples to this sub-space ← dimension reduction
    - $x = [a_1, a_2, a_3, a_4, a_5, a_6, a_7, a_8, \dots, a_m, ]$
    - $x^{sub} = b_1 e_1 + b_2 e_2 + b_3 e_3 + \dots b_n e_n$ 
      - new coordinate in the new subspace is  $[b_n, b_n, b_n, \dots, b_n]$
      - $n < m$

# The Largest Two Principle Component



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  - Gaussian contours
- **PCA face recognition**

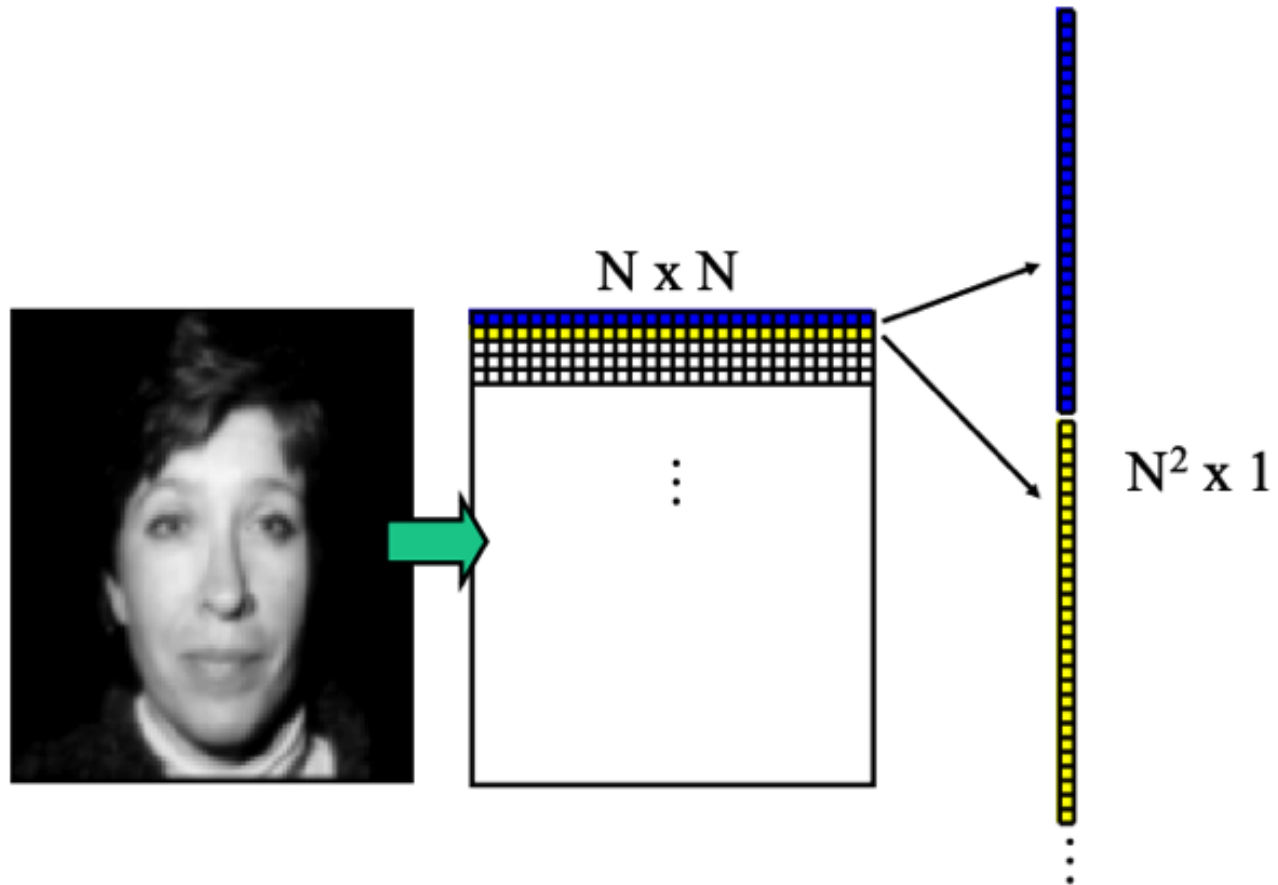
# Face Recognition

- Old method
- Project face images to Gaussian feature space spanning significant variation among known face image
- Significant features in eigenspace for projection are call “EigenFace”
  - The eigenvectors capturing the majority of “variance” of data
    - Largest vector correspond to the largest component variance
  - Project face image (vector) onto selected top eigenvectors

# Face Recognition

- Recognition achieved by comparing weights/coefficients of new face (after projection onto eigenspace) to other stored face weights/coefficients
  - Distance calculation more compact and efficient (uses small number of weights/ coefficients)
- Calculate distance from face space or individual
  - Does it look like a face?
  - Does it look like “Casey”?

# Input Data



Rasterize the image into vector



# Input Data

- Compute mean face image  $\Psi$ 
  - From set of rasterized face images  $\Upsilon_i$  (training image)
- Subtract mean from images
  - Remove the mean using  $\Phi_i = \Upsilon_i - \Psi$
  - From matrix of training faces
    - $A = [\Phi_0 \ \Phi_1 \ \Phi_2 \ \Phi_3 \ \dots \ \Phi_{M-1}]$
    - Matrix size:  $N^2 * M$
- Compute covariance matrix of A

# Eigen Trick

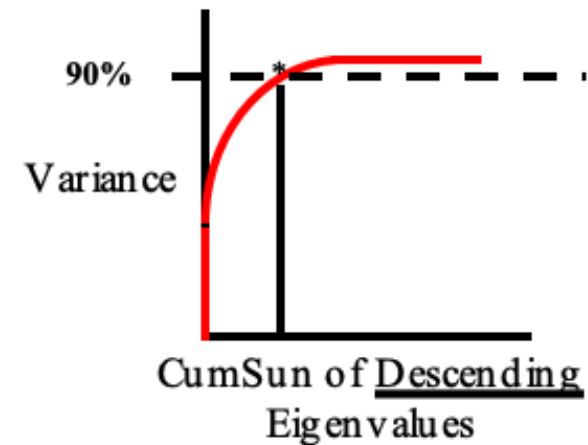
- Compute eigenvectors and eigenvalues of A from its covariance matrix
  - Gaussian sub-space
- Number of face image(M) is much less than the dimension of space ( $N^2$ )
- Thus only M-1 meaningful eigenvectors in the matrix A
  - Remaining eigenvectors have eigenvalues = 0
- Can solve using a much smaller  $M \times M$  matrix and convert back to  $N^2 \times 1$

# Eigen Trick

- Consider the eigenvectors  $v_i$  of  $A^T A$ 
  - Recall  $AA^T$  is how to compute covariance matrix  $K$  for  $\Phi_i$  data
    - $(A^T A) v_i = \lambda_i v_i$
  - Pre-multiplying both sides by  $A$ , we get
    - $A(A^T A) v_i = \lambda_i A v_i$
  - Thus  $u_i = A v_i$  are eigenvectors of  $AA^T$ 
    - Covariance matrix :  $(AA^T)[A v_i] = \lambda_i [A v_i]$
  - Size comparison
    - Matrix  $A^T A$  is size  $M \times M$
    - Matrix  $AA^T$  (a covariance) is size  $N^2 \times N^2$
  - Hence, compute eigenvectors/eigenvalues from  $A^T A$  and use  $u_i = A v_i$  to recover the desired dimensionality
    - Make sure to normalize the  $u_i$  to make them unit vectors

# Compute Eigenvectors

- Retain only top  $m$  eigenvectors ( $u_k$ )
  - Determine from strength of eigenvalues
  - These eigenvectors are the “EigenFaces”
- Accumulate eigenvalues until reach desired percentage of total sum of eigenvalues
  - Pre-sort eigenvalues from largest to smallest
  - Considered as “% variance captured”
  - Typically use around 90%



# Projection into Face Space

- Project each rasterized (mean-subtracted) face image into sub-space (m eigenvectors)
  - Project onto each eigenvector (“EigenFace”)
    - $\omega_T = u_k^T \cdot \Phi_i$
- Keep projection coefficients as representation of rasterized image
  - $\Phi_i \rightarrow \Omega_i = [\omega_1 \omega_2 \omega_3 \dots \omega_m]^T$

# Reconstruction from Face Space

- Reconstruct face image from sub-space
  - Reconstruct from projection coefficients on each eigenvector
  - $\Phi_{recon} = \sum \omega_k u_k$
- Add back mean face
  - $\Upsilon_{recon} = \Phi_{recon} + \Psi$

# Recognition Pipeline

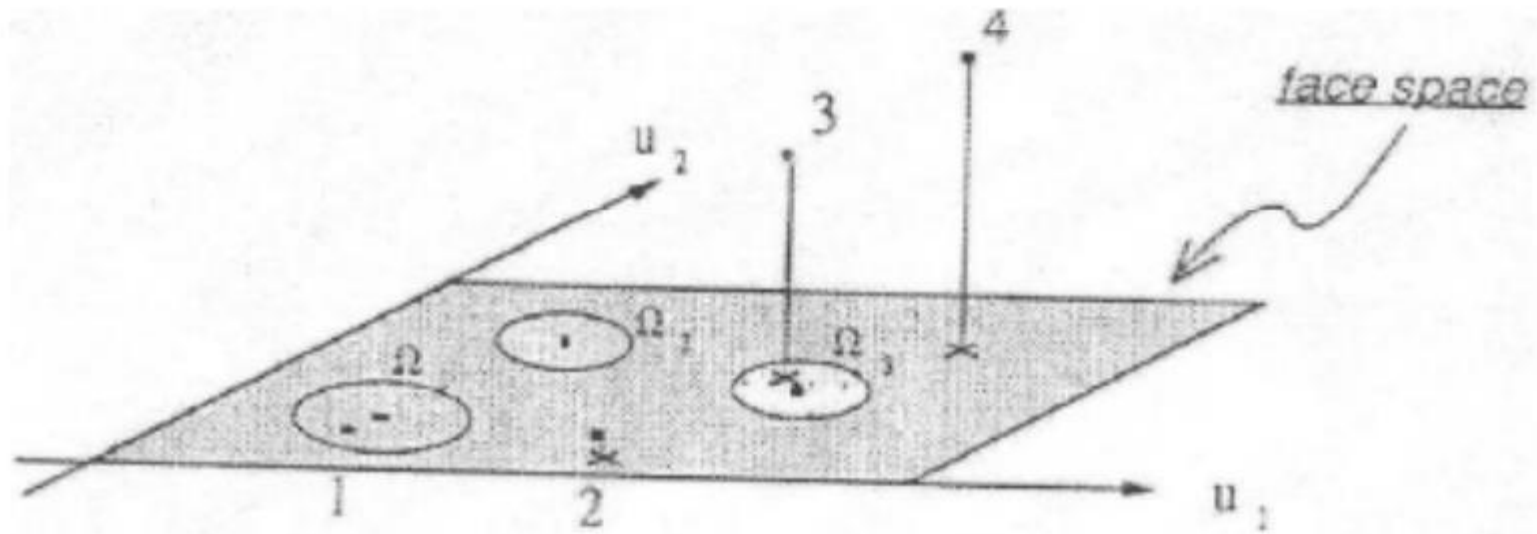
- Get new face image  $\Phi$ 
  - Rasterize
  - Mean-subtract using  $\Psi$
- Compute  $\Omega$
- Use Sum-of-Squared-Error(SSE) to other faces in the database
  - Find best match for database items  $\Omega_i$  (assumes that new face is in the database)
  - $ID = \operatorname{argmin}_i \|\Omega - \Omega_i\|^2$
  - If  $\|\Omega - \Omega_{ID}\|^2 < \textit{Threshold}_{recog}$ ,  $\Omega$  is ID

# Face Detection

- Get new image  $\Phi$ 
  - Rasterize
  - Mean-subtract using  $\Psi$
- Compute  $\Omega$
- Reconstruct image  $\Phi_{recon}$  from  $\Omega$
- Compute  $\|\Phi - \Phi_{recon}\|^2$
- If  $\|\Phi - \Phi_{recon}\|^2 < Threshold_{detect}$ ,  $\Phi$  is face

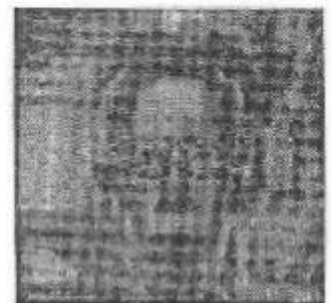
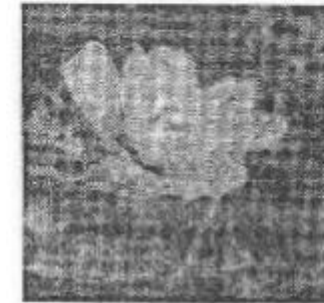


# Face Detection



Input Image

Reconstructed Image



# Principal component (eigenvector) $u_k$



$$\mu + 3\sigma_k u_k$$



$$\mu - 3\sigma_k u_k$$



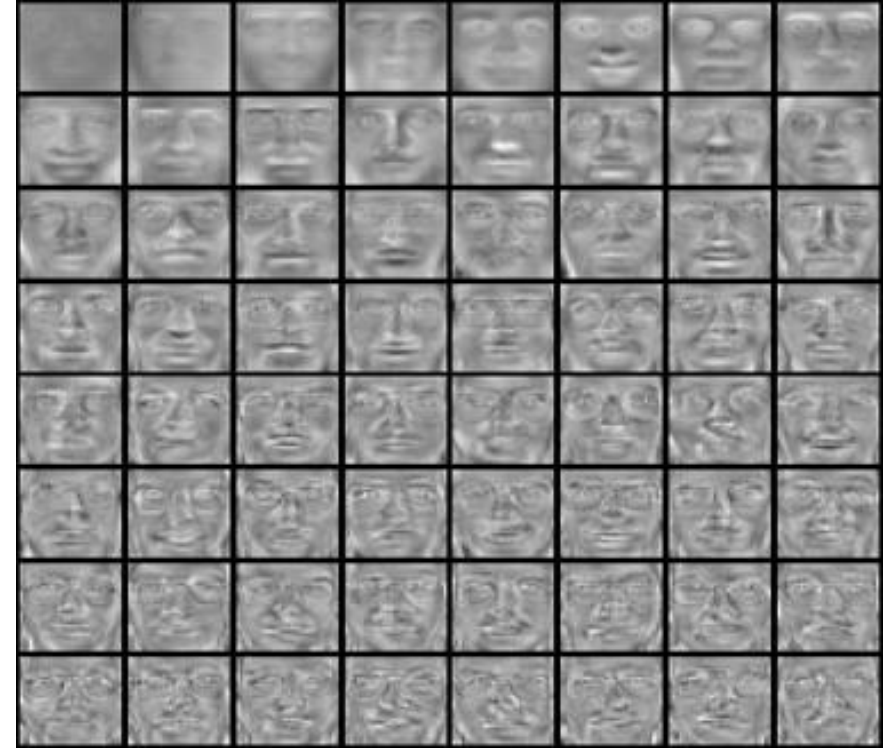
Training dataset



Mean face  $\Psi$



Eigen Faces



Face reconstruction

Query image



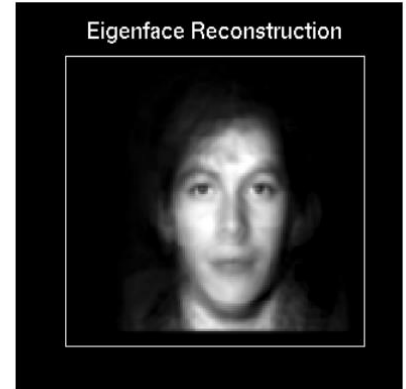
Found image



Input Image



Eigenface Reconstruction



# Limitation

- Background variation and misalignment

