Lecture:

Face Recognition and Feature Reduction

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Recap - Curse of dimensionality

- Assume 5000 points uniformly distributed in the unit hypercube and we want to apply 5-NN. Suppose our query point is at the origin.
	- In 1-dimension, we must go a distance of 5/5000=0.001 on the average to capture 5 nearest neighbors.
	- $-$ In 2 dimensions, we must go $\sqrt{0.001}$ to get a square that contains 0.001 of the volume.
	- $-$ In d dimensions, we must go $\,0.001\right)^{\!1/d}$

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What we will learn today

- Singular value decomposition
- Principal Component Analysis (PCA)
- Image compression

What we will learn today

- Singular value decomposition
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- Image compression

- There are several computer algorithms that can "factorize" a matrix, representing it as the product of some other matrices
- The most useful of these is the Singular Value Decomposition.
- Represents any matrix **A** as a product of three matrices: **UΣVT**
- Python command:

 $-$ [U,S,V]= numpy.linalg.svd(A)

$U\Sigma V^{T} = A$

• Where **U** and **V** are rotation matrices, and **Σ** is a scaling matrix. For example:

$$
\begin{bmatrix} U & \Sigma & V^T & A \\ -.40 & .916 & .40 \end{bmatrix} \times \begin{bmatrix} 5.39 & 0 \\ 0 & .3154 \end{bmatrix} \times \begin{bmatrix} -.05 & .999 \\ .999 & .05 \end{bmatrix} = \begin{bmatrix} 3 & -2 \\ 1 & 5 \end{bmatrix}
$$

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- Beyond 2x2 matrices:
	- $-$ In general, if **A** is *m* x *n*, then **U** will be *m* x *m*, **Σ** will be *m* x *n*, and V^T will be *n* x *n*.
	- $-$ (Note the dimensions work out to produce *m* x *n* after multiplication)

$$
\begin{bmatrix} U & \Sigma \\ -.39 & -.92 \\ -.92 & .39 \end{bmatrix} \times \begin{bmatrix} 9.51 & 0 & 0 \\ 0 & .77 & 0 \end{bmatrix} \times \begin{bmatrix} -.42 & -.57 & -.70 \\ .81 & .11 & -.58 \\ .41 & -.82 & .41 \end{bmatrix} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix}
$$

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- **U** and **V** are always rotation matrices.
	- $-$ Geometric rotation may not be an applicable concept, depending on the matrix. So we call them "unitary" $m \text{airices}$ – each column is a unit vector.
- Σ is a diagonal matrix
	- $-$ The number of nonzero entries $=$ rank of **A**
	- $-$ The algorithm always sorts the entries high to low

$$
\begin{bmatrix} U & \Sigma \\ -.39 & -.92 \\ -.92 & .39 \end{bmatrix} \times \begin{bmatrix} 9.51 & 0 & 0 \\ 0 & .77 & 0 \end{bmatrix} \times \begin{bmatrix} -0.42 & -0.57 & -0.70 \\ 0.81 & 0.11 & -0.58 \\ 0.41 & -.82 & .41 \end{bmatrix} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix}
$$

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- We've discussed SVD in terms of geometric transformation matrices
- But SVD of an image matrix can also be very useful
- To understand this, we'll look at a less geometric interpretation of what SVD is doing

$$
\begin{bmatrix} U & \Sigma \\ -.39 & -.92 \\ -.92 & .39 \end{bmatrix} \times \begin{bmatrix} 9.51 & 0 & 0 \\ 0 & .77 & 0 \end{bmatrix} \times \begin{bmatrix} -.42 & -.57 & -.70 \\ .81 & .11 & -.58 \\ .41 & -.82 & .41 \end{bmatrix} = \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix}
$$

- Look at how the multiplication works out, left to right:
- Column 1 of **U** gets scaled by the first value from **Σ**.

• The resulting vector gets scaled by row 1 of V^T to produce a contribution to the columns of A

• Each product of (*column i of U*)**·**(*value i from Σ*)**·**(*row i of* V^T) produces a component of the final A.

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- We're building **A** as a linear combination of the columns of *U*
- Using all columns of U, we'll rebuild the original matrix perfectly
- But, in real-world data, often we can just use the first few columns of *U* and we'll get something close (e.g. the first $A_{partial}$, above)

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- We can call those first few columns of U the *Principal Components* of the data
- They show the major patterns that can be added to produce the columns of the original matrix
- The rows of **V^T** show how the *principal components* are mixed to produce the columns of the matrix

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We can look at **Σ** to see that the first column has a large effect

while the second column has a much smaller effect in this example

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- For this image, using only the first 10 of 300 principal components produces a recognizable reconstruction
- So, SVD can be used for image compression

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SVD for symmetric matrices

• If A is a symmetric matrix, it can be decomposed as the following:

$$
A=\Phi\Sigma\Phi^T
$$

• Compared to a traditional SVD decomposition, $U = V^T$ and is an orthogonal matrix.

Principal Component Analysis

- Remember, columns of U are the *Principal Components* of the data: the major patterns that can be added to produce the columns of the original matrix
- One use of this is to construct a matrix where each column is a separate data sample
- Run SVD on that matrix, and look at the first few columns of **U** to see patterns that are common among the columns
- This is called *Principal Component Analysis* (or PCA) of the data samples

Principal Component Analysis

- Often, raw data samples have a lot of redundancy and patterns
- PCA can allow you to represent data samples as weights on the principal components, rather than using the original raw form of the data
- By representing each sample as just those weights, you can represent just the "meat" of what's different between samples.
- This minimal representation makes machine learning and other algorithms much more efficient

How is SVD computed?

- For this class: tell PYTHON to do it. Use the result.
- But, if you're interested, one computer algorithm to do it makes use of Eigenvectors!

Eigenvector definition

- Suppose we have a square matrix **A**. We can solve for vector x and scalar λ such that $Ax = \lambda x$
- In other words, find vectors where, if we transform them with A, the only effect is to scale them with no change in direction.
- These vectors are called eigenvectors (German for "self vector" of the matrix), and the scaling factors λ are called eigenvalues
- An *m* x *m* matrix will have ≤ *m* eigenvectors where λ is nonzero

Finding eigenvectors

• Computers can find an x such that $Ax = \lambda x$ using this iterative algorithm:

 $- X =$ random unit vector

- while(x hasn't converged)
	- $X = Ax$
	- normalize x
- x will quickly converge to an eigenvector
- Some simple modifications will let this algorithm find all eigenvectors

Finding SVD

- Eigenvectors are for square matrices, but SVD is for all matrices
- To do svd(A), computers can do this:
	- $-$ Take eigenvectors of AA^T (matrix is always square).
		- These eigenvectors are the columns of **U**.
		- Square root of eigenvalues are the singular values (the entries of **Σ**).
	- $-$ Take eigenvectors of ATA (matrix is always square).
		- These eigenvectors are columns of **V** (or rows of **V**^T)

Finding SVD

- Moral of the story: SVD is fast, even for large matrices
- It's useful for a lot of stuff
- There are also other algorithms to compute SVD or part of the SVD
	- Python's np.linalg.svd() command has options to efficiently compute only what you need, if performance becomes an issue

A detailed geometric explanation of SVD is here: http://www.ams.org/samplings/feature-column/fcarc-svd

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What we will learn today

- Introduction to face recognition
- Principal Component Analysis (PCA)
- Image compression

Covariance

- Variance and Covariance are a measure of the "spread" of a set of points around their center of mass (mean)
- Variance measure of the deviation from the mean for points in one dimension e.g. heights
- Covariance as a measure of how much each of the dimensions vary from the mean with respect to each other.
- Covariance is measured between 2 dimensions to see if there is a relationship between the 2 dimensions e.g. number of hours studied & marks obtained.
- The covariance between one dimension and itself is the variance

Covariance

$$
covariance (X,Y) = \frac{\sum_{i=1}^{n} (X_i - X) (Y_i - Y)}{(n-1)}
$$

• So, if you had a 3-dimensional data set (x,y,z), then you could measure the covariance between the x and y dimensions, the y and z dimensions, and the x and z dimensions. Measuring the covariance between x and x, or y and y, or z and z would give you the variance of the x, y and z dimensions respectively

Covariance matrix

• Representing Covariance between dimensions as a matrix e.g. for 3 dimensions

$$
C = \begin{bmatrix} \text{cov}(x,x) - \text{cov}(x,y) & \text{cov}(x,z) \\ \text{cov}(y,x) - \text{cov}(y,y) & \text{cov}(x,z) \\ \text{cov}(z,x) & \text{cov}(z,y) - \text{cov}(z,z) \end{bmatrix}
$$

 Variances

- Diagonal is the variances of x, y and z
- $cov(x,y) = cov(y,x)$ hence matrix is symmetrical about the diagonal
- N-dimensional data will result in NxN covariance matrix

Covariance

- What is the interpretation of covariance calculations?
	- e.g.: 2 dimensional data set
	- $-x$: number of hours studied for a subject
	- $-$ y: marks obtained in that subject
	- covariance value is say: 104.53
	- $-$ what does this value mean?

Covariance interpretation

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Covariance interpretation

- Exact value is not as important as it's sign.
- A positive value of covariance indicates both dimensions increase or decrease together e.g. as the number of hours studied increases, the marks in that subject increase.
- A negative value indicates while one increases the other decreases, or vice-versa e.g. active social life at PSU vs performance in CS dept.
- If **covariance is zero**: the two dimensions are independent of each other e.g. heights of students vs the marks obtained in a subject

Example data

Covariance between the two axis is high. Can we reduce the number of dimensions to just 1?

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Geometric interpretation of PCA

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Geometric interpretation of PCA

• Let's say we have a set of 2D data points x. But we see that all the points lie on a line in 2D.

• So, 2 dimensions are redundant to express the data. We can express all the points with just one dimension.

PCA: Principle Component Analysis

- Given a set of points, how do we know if they can be compressed like in the previous example?
	- $-$ The answer is to look into the correlation between the points
	- $-$ The tool for doing this is called PCA

PCA Formulation

- Basic idea:
	- $-$ If the data lives in a subspace, it is going to look very flat when viewed from the full space, e.g.

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PCA Formulation

- Assume x is Gaussian with covariance Σ.
- Recall that a gaussian is defined with it's mean and variance:

 $\mathbf{X} \sim \mathcal{N}(\boldsymbol{\mu}, \boldsymbol{\Sigma})$

• Recall that μ and Σ of a gaussian are defined as:

$$
\boldsymbol{\mu} = \mathrm{E}[\mathbf{X}] = [\mathrm{E}[X_1], \mathrm{E}[X_2], \ldots, \mathrm{E}[X_k]]^\mathrm{T}
$$

 $\mathbf{\Sigma} =: \mathrm{E}[(\mathbf{X}-\boldsymbol{\mu})(\mathbf{X}-\boldsymbol{\mu})^{\mathrm{T}}] = [\mathrm{Cov}[X_i,X_j]; 1\leq i,j\leq k]$

 X_2'

 $\dot{\varphi}_2$

 λ_2 λ_1

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 X_1

 Φ_1

PCA formulation

• Since gaussians are symmetric, it's covariance matrix is also a symmetric matrix. So we can express it as:

$$
-\Sigma = U\Lambda U^{T} = U\Lambda^{1/2}(U\Lambda^{1/2})^{T}
$$

$$
\mathbf{X}\ \sim \mathcal{N}(\boldsymbol{\mu},\mathbf{\Sigma})\iff \mathbf{X}\ \sim \boldsymbol{\mu}+\mathbf{U}\boldsymbol{\Lambda}^{1/2}\mathcal{N}(0,\mathbf{I})
$$

$$
\iff \mathbf{X}\ \sim\boldsymbol{\mu}+\mathbf{U}\mathcal{N}(0,\boldsymbol{\Lambda}).
$$

PCA Formulation

• If x is Gaussian with covariance Σ ,

- $-$ Principal components ϕ_i are the eigenvectors of Σ
- $-$ Principal lengths λ_i are the eigenvalues of Σ

- by computing the eigenvalues we know the data is
	- Not flat if λ_1 ≈ λ_2
	- Flat if λ_1 >> λ_2

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PCA Algorithm (training)

Given sample
$$
\mathcal{D} = {\mathbf{x}_1, \dots, \mathbf{x}_n}
$$
, $x_i \in \mathcal{R}^d$

• compute sample mean: $\hat{\mu} = \frac{1}{n} \sum_i (\mathbf{x}_i)$

• compute sample covariance: $\hat{\Sigma} = \frac{1}{n} \sum_i (\mathbf{x}_i - \hat{\mu}) (\mathbf{x}_i - \hat{\mu})^T$

• compute eigenvalues and eigenvectors of $\hat{\Sigma}$

$$
\hat{\Sigma} = \Phi \Lambda \Phi^T, \ \Lambda = diag(\sigma_1^2, \dots, \sigma_n^2) \ \Phi^T \Phi = I
$$

• order eigenvalues
$$
\sigma_1^2 > ... > \sigma_n^2
$$

• if, for a certain k, $\sigma_k << \sigma_1$ eliminate the eigenvalues and eigenvectors above k .

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PCA Algorithm (testing)

• Given principal compoenents $\phi_i, i \in 1, \ldots, k$ and a test sample $\mathcal{T} = \{\mathbf{t}_1, \ldots, \mathbf{t}_n\}, \ \ t_i \in \mathcal{R}^d$

• subtract mean to each point $t'_i = t_i - \hat{\mu}$

• project onto eigenvector space $y_i = At'_i$ where

$$
\mathbf{A} = \begin{bmatrix} \phi_1^T \\ \vdots \\ \phi_k^T \end{bmatrix}
$$

• use $\mathcal{T}' = \{y_1, \ldots, y_n\}$ to estimate class conditional densities and do all further processing on y.

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PCA by SVD

- An alternative manner to compute the principal components, based on singular value decomposition
- Quick reminder: SVD
	- $-$ Any real n x m matrix (n>m) can be decomposed as

- $-$ Where M is an (n x m) column orthonormal matrix of left singular vectors (columns of M)
- $-$ T is an (m x m) diagonal matrix of singular values
- $-$ N^T is an (m x m) row orthonormal matrix of right singular vectors (columns of N)

$$
M^T M = I \qquad N^T N = I
$$

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PCA by SVD

• To relate this to PCA, we consider the data matrix

• The sample mean is

$$
\mu = \frac{1}{n} \sum_{i} X_{i} = \frac{1}{n} \begin{bmatrix} 1 & 1 \\ X_{1} & \cdots & X_{n} \\ 1 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix} = \frac{1}{n} X 1
$$

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PCA by SVD

- Center the data by subtracting the mean to each column of X
- The centered data matrix is

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PCA by SVD

• The sample covariance matrix is

$$
\Sigma = \frac{1}{n} \sum_{i} (x_i - \mu)(x_i - \mu)^T = \frac{1}{n} \sum_{i} x_i^c (x_i^c)^T
$$

where $\mathsf{x_i}^\mathsf{c}$ is the ith column of X_c

• This can be written as

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PCA by SVD

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• The matrix

$$
\mathbf{X}_c^T = \begin{bmatrix} - & \mathbf{X}_1^c & - \\ \vdots & & \\ - & \mathbf{X}_n^c & - \end{bmatrix}
$$

 $=$

is real (n x d). Assuming n>d it has SVD decomposition

$$
\boldsymbol{X}_c^T = \boldsymbol{\mathrm{M}\Pi\boldsymbol{\mathrm{N}}^T} \qquad \qquad \boldsymbol{\mathrm{M}^T\boldsymbol{\mathrm{M}}} = \boldsymbol{I} \qquad \qquad \boldsymbol{\mathrm{N}^T\boldsymbol{\mathrm{N}}} = \boldsymbol{I}
$$

and

$$
\Sigma = \frac{1}{n} X_c X_c^T = \frac{1}{n} \text{NIM}^T \text{MIN}^T = \frac{1}{n} \text{NII}^2 \text{N}^T
$$

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PCA by SVD

- Note that N is (d x d) and orthonormal, and Π^2 is diagonal. This is just the eigenvalue decomposition of Σ
- It follows that
	- $-$ The eigenvectors of Σ are the columns of N
	- $-$ The eigenvalues of Σ are

$$
\lambda_i = \frac{1}{n} \pi_i^2
$$

This gives an alternative algorithm for PCA

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PCA by SVD

- In summary, computation of PCA by SVD
- Given X with one example per column
	- $-$ Create the centered data matrix

$$
X_c^T = \left(I - \frac{1}{n} 11^T\right) X^T
$$

Compute its SVD

$$
X_c^T = \text{MIN}^T
$$

Principal components are columns of N, eigenvalues are

$$
\lambda_i = \frac{1}{n} \pi_i^2
$$

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Rule of thumb for finding the number of PCA components

- A natural measure is to pick the eigenvectors that explain p% of the data variability
	- Can be done by plotting the ratio r_k as a function of k

 $-$ E.g. we need 3 eigenvectors to cover 70% of the variability of this dataset

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What we will learn today

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Original Image

- Divide the original 372x492 image into patches:
	- Each patch is an instance that contains 12x12 pixels on a grid
- View each as a 144-D vector

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L_2 error and PCA dim

PCA compression: 144D) 60D

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PCA compression: 144D) 16D

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16 most important eigenvectors

PCA compression: 144D) 6D

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6 most important eigenvectors

PCA compression: 144D) 3D

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3 most important eigenvectors

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PCA compression: 144D) 1D

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