

10-701 Introduction to Machine Learning

Principal Component Analysis and Dimensionality Reduction

Readings:

Bishop Ch. 12

Murphy Ch. 12

Matt Gormley Lecture 14 October 24, 2016

DIMENSIONALITY REDUCTION

Big & High-Dimensional Data

High-Dimensions = Lot of Features

Document classification

Features per document =
thousands of words/unigrams
millions of bigrams, contextual
information



Surveys - Netflix

480189 users x 17770 movies

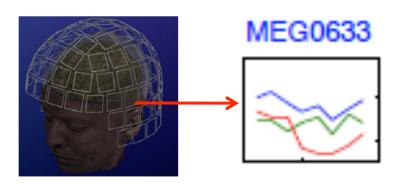
	movie 1	movie 2	movie 3	movie 4	movie 5	movie 6
Tom	5	?	?	1	3	?
George	?	?	3	1	2	5
Susan	4	3	1	?	5	1
Beth	4	3	?	2	4	2

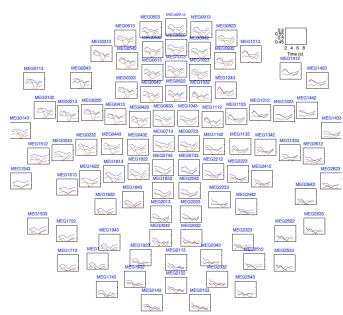
Big & High-Dimensional Data

• High-Dimensions = Lot of Features

MEG Brain Imaging

120 locations x 500 time points x 20 objects





Or any high-dimensional image data



Big & High-Dimensional Data.

 Useful to learn lower dimensional representations of the data.

Learning Representations

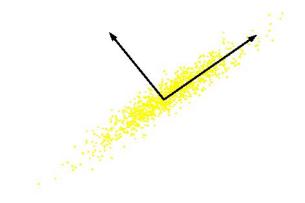
PCA, Kernel PCA, ICA: Powerful unsupervised learning techniques for extracting hidden (potentially lower dimensional) structure from high dimensional datasets.

Useful for:

- Visualization
 - More efficient use of resources (e.g., time, memory, communication)
- Statistical: fewer dimensions → better generalization
- Noise removal (improving data quality)
- Further processing by machine learning algorithms

Principal Component Analysis (PCA)

What is PCA: Unsupervised technique for extracting variance structure from high dimensional datasets.



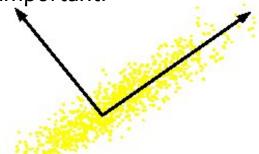
 PCA is an orthogonal projection or transformation of the data into a (possibly lower dimensional) subspace so that the variance of the projected data is maximized.

Principal Component Analysis (PCA)

Intrinsically lower dimensional than the dimension of the ambient space.



If we rotate data, again only one coordinate is more important.

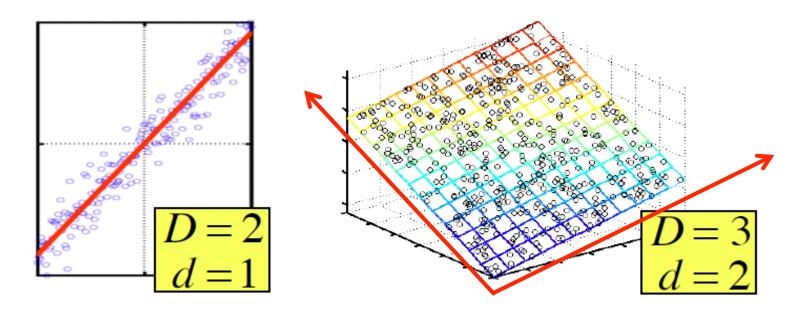


Only one relevant feature

Both features are relevant

Question: Can we transform the features so that we only need to preserve one latent feature?

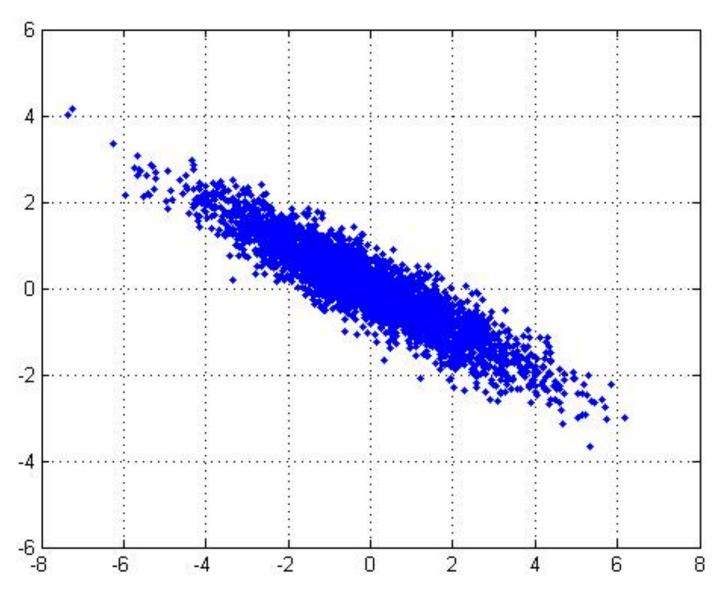
Principal Component Analysis (PCA)



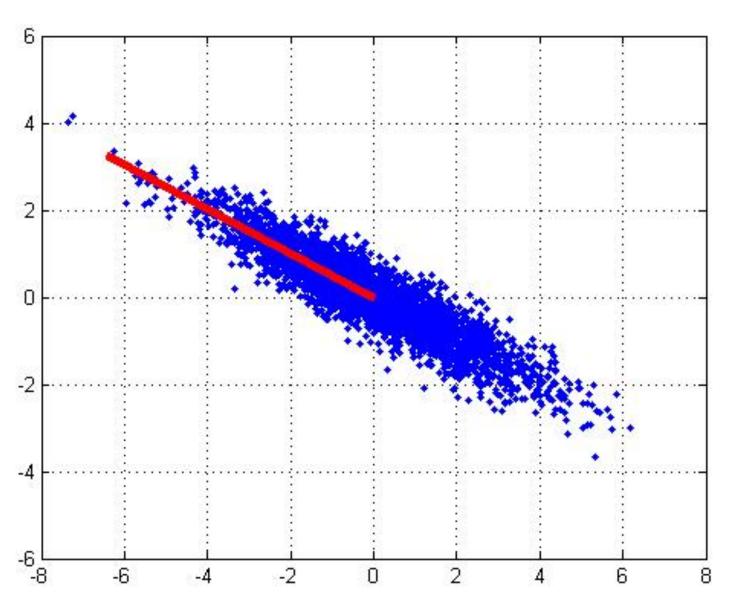
In case where data lies on or near a low d-dimensional linear subspace, axes of this subspace are an effective representation of the data.

Identifying the axes is known as Principal Components Analysis, and can be obtained by using classic matrix computation tools (Eigen or Singular Value Decomposition).

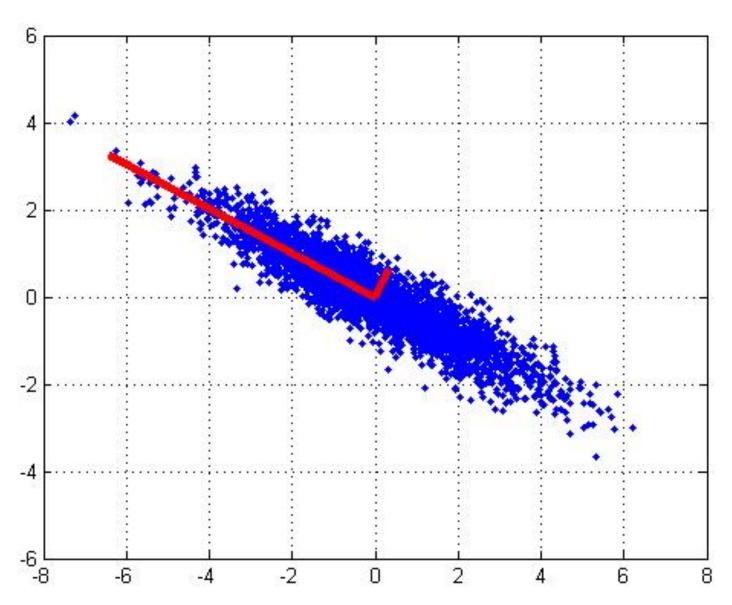
2D Gaussian dataset



1st PCA axis



2nd PCA axis



PCA ALGORITHMS

PCA algorithm I (sequential)

Given the **centered** data $\{x_1, ..., x_m\}$, compute the principal vectors:

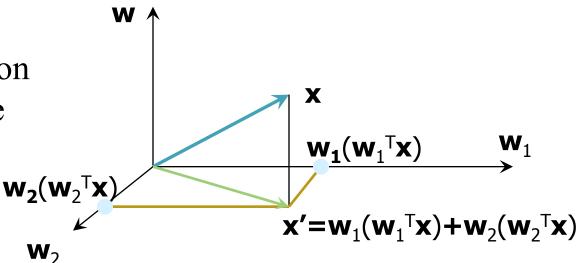
$$\mathbf{w}_1 = \arg\max_{\|\mathbf{w}\|=1} \frac{1}{m} \sum_{i=1}^m \{(\mathbf{w}^T \mathbf{x}_i)^2\} \qquad 1^{\text{st}} \text{ PCA vector}$$

We maximize the variance of projection of \mathbf{x}

$$\mathbf{w}_{k} = \arg \max_{\|\mathbf{w}\|=1} \frac{1}{m} \sum_{i=1}^{m} \{ [\mathbf{w}^{T} (\mathbf{x}_{i} - \sum_{j=1}^{k-1} \mathbf{w}_{j} \mathbf{w}_{j}^{T} \mathbf{x}_{i})]^{2} \}$$

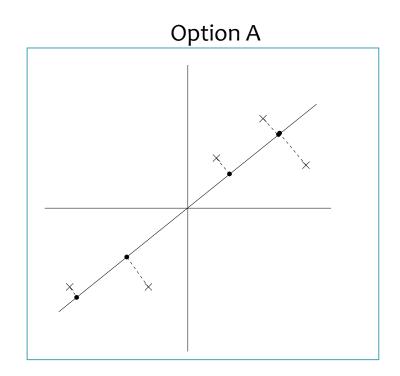
$$\mathbf{x'} \text{ PCA reconstruction}$$

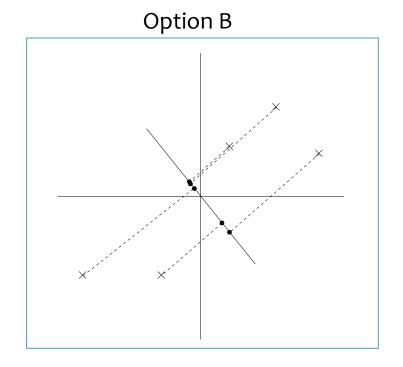
We maximize the variance of the projection in the residual subspace



Maximizing the Variance

- Consider the two projections below
- Which maximizes the variance?





PCA algorithm I (sequential)

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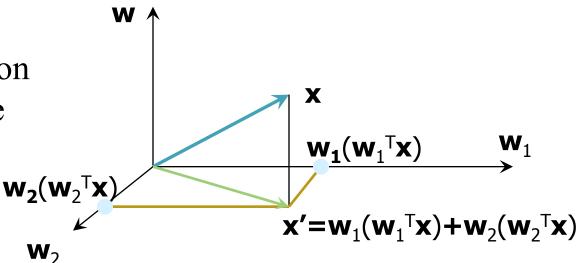
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We maximize the variance of the projection in the residual subspace



PCA algorithm II (sample covariance matrix)

• Given data $\{x_1, ..., x_m\}$, compute covariance matrix Σ

$$\Sigma = \frac{1}{m} \sum_{i=1}^{m} (\mathbf{x}_i - \overline{\mathbf{x}}) (\mathbf{x} - \overline{\mathbf{x}})^T \quad \text{where} \quad \overline{\mathbf{x}} = \frac{1}{m} \sum_{i=1}^{m} \mathbf{x}_i$$

$$\overline{\mathbf{x}} = \frac{1}{m} \sum_{i=1}^{m} \mathbf{x}_{i}$$

• **PCA** basis vectors = the eigenvectors of Σ

We get the eigvectors using an eigendecomposition. Power iteration (Von Mises iteration is a standard algorithm for this)

Larger eigenvalue ⇒ more important eigenvectors





Maximise $\mathbf{u}^{\mathsf{T}}\mathbf{X}\mathbf{X}^{\mathsf{T}}\mathbf{u}$

s.t $\mathbf{u}^{\mathsf{T}}\mathbf{u} = 1$

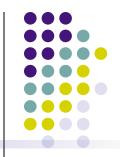
Construct Langrangian $\mathbf{u}^{\mathsf{T}}\mathbf{X}\mathbf{X}^{\mathsf{T}}\mathbf{u} - \lambda \mathbf{u}^{\mathsf{T}}\mathbf{u}$

Vector of partial derivatives set to zero

$$\mathbf{x}\mathbf{x}^{\mathsf{T}}\mathbf{u} - \lambda\mathbf{u} = (\mathbf{x}\mathbf{x}^{\mathsf{T}} - \lambda\mathbf{I})\mathbf{u} = 0$$

As $\mathbf{u} \neq \mathbf{0}$ then \mathbf{u} must be an eigenvector of $\mathbf{X}\mathbf{X}^{\mathsf{T}}$ with eigenvalue λ





 For symmetric matrices, eigenvectors for distinct eigenvalues are orthogonal

$$Sv_{\{1,2\}} = \lambda_{\{1,2\}}v_{\{1,2\}}$$
, and $\lambda_1 \neq \lambda_2 \Rightarrow v_1 \cdot v_2 = 0$

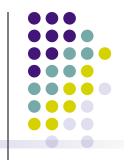
All eigenvalues of a real symmetric matrix are real.

if
$$|S - \lambda I| = 0$$
 and $S = S^T \Rightarrow \lambda \in \Re$

 All eigenvalues of a positive semidefinite matrix are nonnegative

$$\forall w \in \Re^n, w^T Sw \ge 0$$
, then if $Sv = \lambda v \Rightarrow \lambda \ge 0$





• Let $S \in \mathbb{R}^{m \times m}$ be a square matrix with m linearly independent eigenvectors (a "non-defective" matrix)

• Theorem: Exists an eigen decomposition

distinct eigenvalues

$$\mathbf{S} = \mathbf{U} \boldsymbol{\Lambda} \mathbf{U}^{-1}$$
 diagonal

(cf. matrix diagonalization theorem)

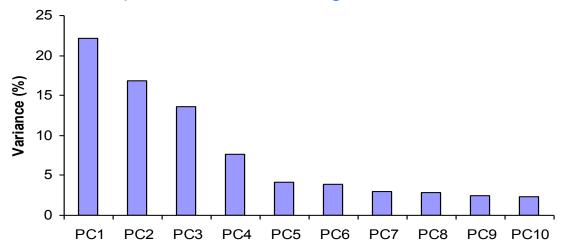
- Columns of *U* are eigenvectors of *S*
- ullet Diagonal elements of $oldsymbol{\Lambda}$ are eigenvalues of $oldsymbol{S}$

$$\Lambda = \operatorname{diag}(\lambda_1, \dots, \lambda_m), \ \lambda_i \ge \lambda_{i+1}$$





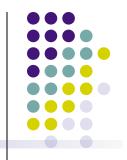
- For n original dimensions, sample covariance matrix is nxn, and has up to n eigenvectors. So n PCs.
- Where does dimensionality reduction come from?
 Can *ignore* the components of lesser significance.



You do lose some information, but if the eigenvalues are small, you don't lose much

- n dimensions in original data
- calculate n eigenvectors and eigenvalues
- choose only the first p eigenvectors, based on their eigenvalues
- final data set has only p dimensions





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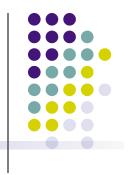
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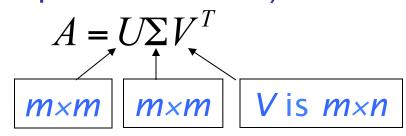
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For an $m \times n$ matrix \mathbf{A} of rank r there exists a factorization (Singular Value Decomposition = \mathbf{SVD}) as follows:



The columns of U are orthogonal eigenvectors of AA^T .

The columns of V are orthogonal eigenvectors of A^TA .

Eigenvalues $\lambda_1 \dots \lambda_r$ of AA^T are the eigenvalues of A^TA .

$$\sigma_{i} = \sqrt{\lambda_{i}}$$

$$\Sigma = diag(\sigma_{1}...\sigma_{r})$$
Singular values.

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PCA: Two Interpretations

E.g., for the first component.

Maximum Variance Direction: 1st PC a vector v such that projection on to this vector capture maximum variance in the data (out of all possible one dimensional projections)

$$\frac{1}{n} \sum_{i=1}^{n} (\mathbf{v}^T \mathbf{x}_i)^2 = \mathbf{v}^T \mathbf{X} \mathbf{X}^T \mathbf{v}$$

Minimum Reconstruction Error: 1st PC a vector v such that projection on to this vector yields minimum MSE reconstruction

$$\frac{1}{n} \sum_{i=1}^{n} \|\mathbf{x}_i - (\mathbf{v}^T \mathbf{x}_i) \mathbf{v}\|^2$$

PCA: Two Interpretations

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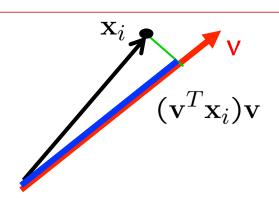
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black² is fixed (it's just the data)

So, maximizing blue² is equivalent to minimizing green²



PCA algorithm III (SVD of the data matrix)

Singular Value Decomposition of the centered data matrix X.

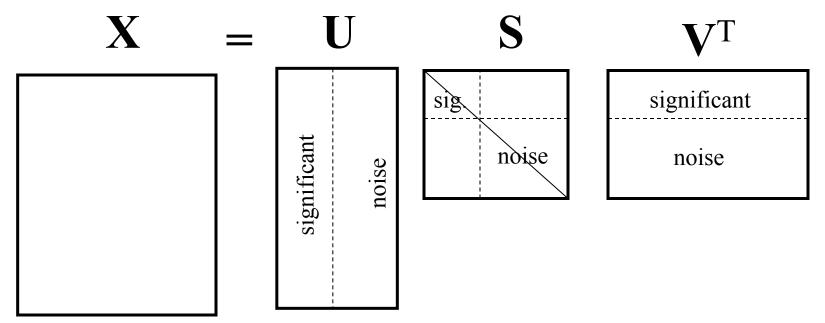
$$\mathbf{X} = [\mathbf{x}_1, \dots, \mathbf{x}_m] \in \mathbb{R}^{N \times m}$$
,

samples

m: number of instances,

N: dimension

$$\mathbf{X}_{\text{features} \times \text{samples}} = \mathbf{U} \mathbf{S} \mathbf{V}^{\text{T}}$$



Slide from Barnabas Poczos

PCA algorithm III

Columns of U

- the principal vectors, { $\mathbf{u}^{(1)}$, ..., $\mathbf{u}^{(k)}$ }
- orthogonal and has unit norm so $U^{T}U = I$
- Can reconstruct the data using linear combinations of { u⁽¹⁾, ..., u^(k) }

Matrix S

- Diagonal
- Shows importance of each eigenvector

Columns of V^T

The coefficients for reconstructing the samples

SVD and **PCA**



- The first root is called the prinicipal eigenvalue which has an associated orthonormal (u^Tu = 1) eigenvector u
- Subsequent roots are ordered such that $\lambda_1 > \lambda_2 > ... > \lambda_M$ with rank(**D**) non-zero values.
- Eigenvectors form an orthonormal basis i.e. $\mathbf{u}_i^{\mathsf{T}}\mathbf{u}_j = \delta_{ij}$
- The eigenvalue decomposition of XX^T = UΣU^T
- where $\mathbf{U} = [\mathbf{u}_1, \mathbf{u}_2, ..., \mathbf{u}_M]$ and $\mathbf{\Sigma} = \text{diag}[\lambda_1, \lambda_2, ..., \lambda_M]$
- Similarly the eigenvalue decomposition of X^TX = VΣV^T
- The SVD is closely related to the above X=U Σ^{1/2} V^T
- The left eigenvectors U, right eigenvectors V,
- singular values = square root of eigenvalues.





Solution via SVD

$$A_k = U \operatorname{diag}(\sigma_1, ..., \sigma_k, \underbrace{0, ..., 0}) V^T$$
set smallest r-k
singular values to zero

$$A_k = \sum_{i=1}^k \sigma_i u_i v_i^T - column notation: sum of rank 1 matrices$$

Approximation error



- How good (bad) is this approximation?
- It's the best possible, measured by the Frobenius norm of the error:

$$\min_{X:rank(X)=k} \|A - X\|_F = \|A - A_k\|_F = \sigma_{k+1}$$

where the σ_i are ordered such that $\sigma_i \ge \sigma_{i+1}$.

Suggests why Frobenius error drops as *k* increased.

Slides from Barnabas Poczos

Original sources include:

- Karl Booksh Research group
- Tom Mitchell
- Ron Parr

PCA EXAMPLES

Face recognition

Challenge: Facial Recognition

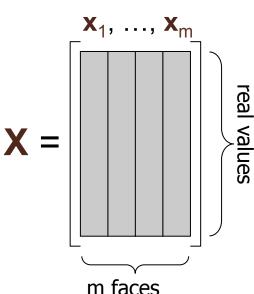
- Want to identify specific person, based on facial image
- Robust to glasses, lighting,...
 - → Can't just use the given 256 x 256 pixels



Applying PCA: Eigenfaces

Method A: Build a PCA subspace for each person and check which subspace can reconstruct the test image the bestMethod B: Build one PCA database for the whole dataset and then classify based on the weights.





- Example data set: Images of faces
 - Famous Eigenface approach
 [Turk & Pentland], [Sirovich & Kirby]
- Each face x is ...
- 256 × 256 values (luminance at location)
 - **x** in $\Re^{256 \times 256}$ (view as 64K dim vector)
- Form $\mathbf{X} = [\mathbf{x}_1, ..., \mathbf{x}_m]$ centered data mtx
 - Compute $\Sigma = XX^T$
 - Problem: Σ is 64K \times 64K ... HUGE!!!

Computational Complexity

- Suppose m instances, each of size N
 - Eigenfaces: m=500 faces, each of size N=64K
- Given N×N covariance matrix Σ, can compute
 - all N eigenvectors/eigenvalues in O(N³)
 - first k eigenvectors/eigenvalues in O(k N²)

But if N=64K, EXPENSIVE!

A Clever Workaround

- Note that m<<64K
- Use L=X^TX instead of Σ=XX^T
- If v is eigenvector of L then Xv is eigenvector of Σ

Proof:
$$\mathbf{L} \ \mathbf{v} = \gamma \ \mathbf{v}$$

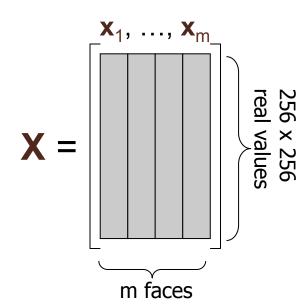
$$\mathbf{X}^{T}\mathbf{X} \ \mathbf{v} = \gamma \ \mathbf{v}$$

$$\mathbf{X} \ (\mathbf{X}^{T}\mathbf{X} \ \mathbf{v}) = \mathbf{X}(\gamma \ \mathbf{v}) = \gamma$$

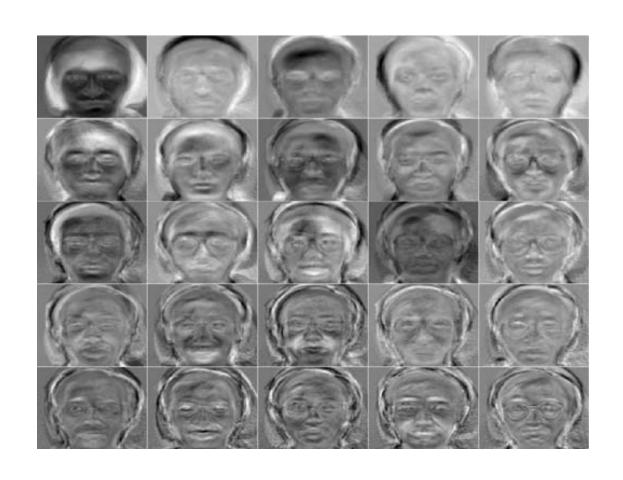
$$\mathbf{X} \mathbf{v}$$

$$(\mathbf{X}\mathbf{X}^{T})\mathbf{X} \ \mathbf{v} = \gamma \ (\mathbf{X}\mathbf{v})$$

$$\mathbf{\Sigma} \ (\mathbf{X}\mathbf{v}) = \gamma \ (\mathbf{X}\mathbf{v})$$



Principle Components (Method B)



Reconstructing... (Method B)



- ... faster if train with...
 - only people w/out glasses
 - same lighting conditions

Shortcomings

- Requires carefully controlled data:
 - All faces centered in frame
 - Same size
 - Some sensitivity to angle
- Alternative:
 - "Learn" one set of PCA vectors for each angle
 - Use the one with lowest error
- Method is completely knowledge free
 - (sometimes this is good!)
 - Doesn't know that faces are wrapped around 3D objects (heads)
 - Makes no effort to preserve class distinctions

Facial expression recognition

Happiness subspace (method A)





















Disgust subspace (method A)













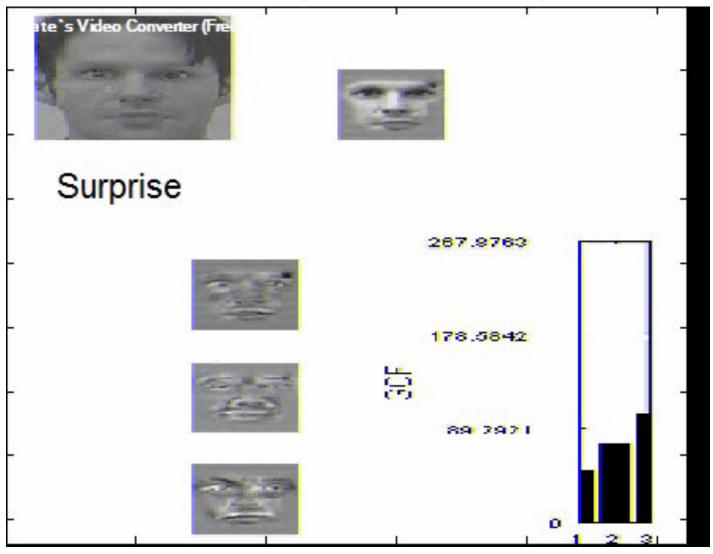




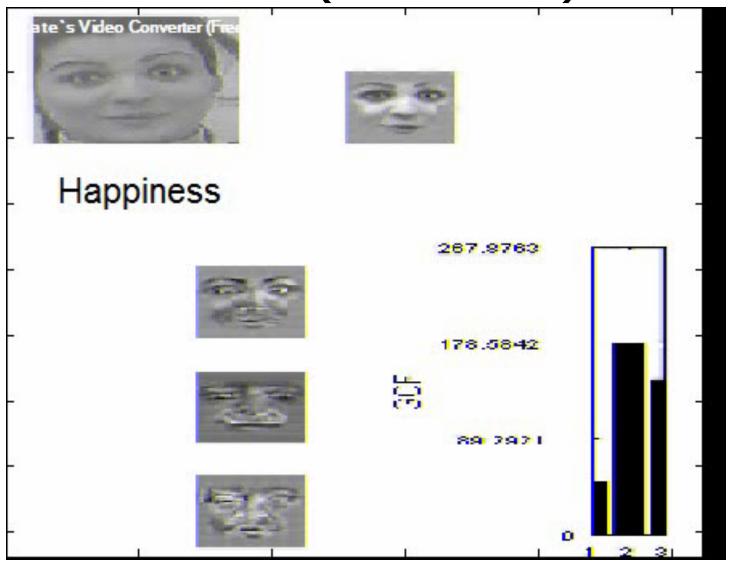




Facial Expression Recognition Movies (method A)



Facial Expression Recognition Movies (method A)



Facial Expression Recognition Movies (method A)

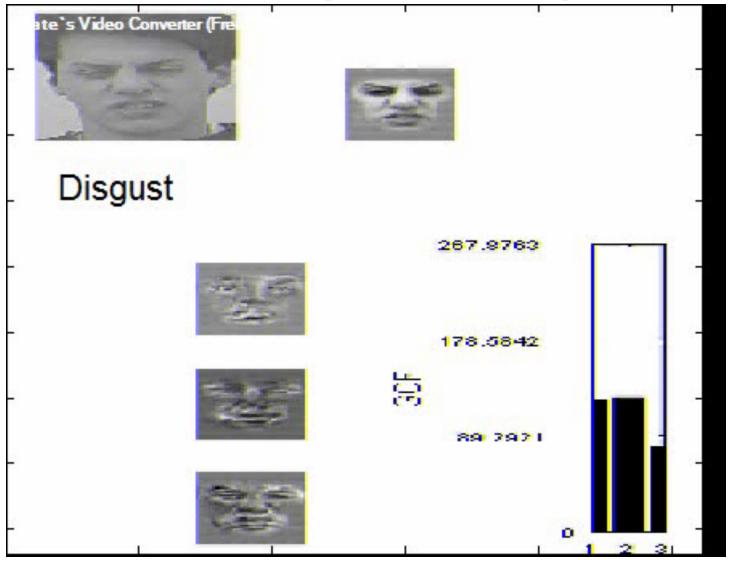


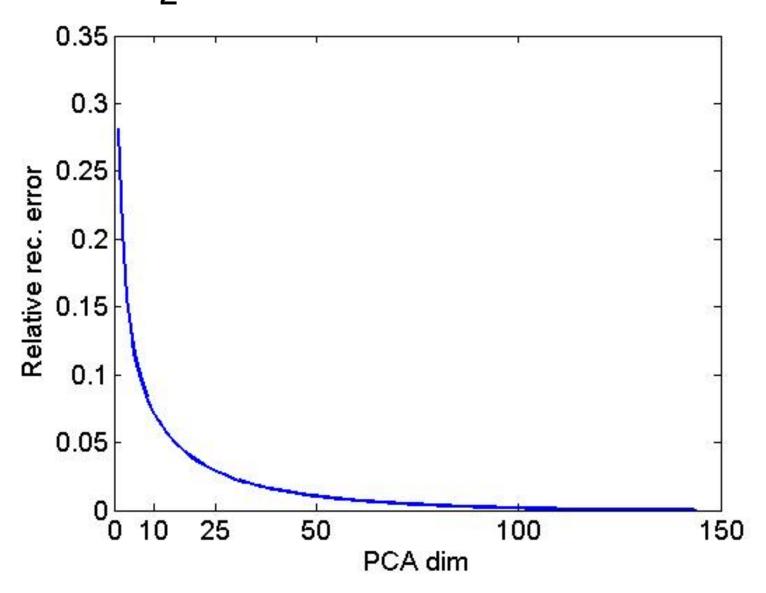
Image Compression

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

L₂ error and PCA dim



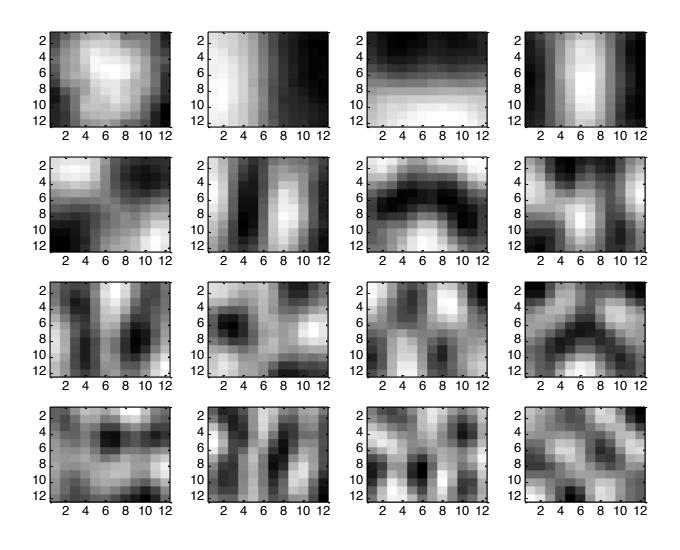
PCA compression: 144D) 60D



PCA compression: 144D) 16D



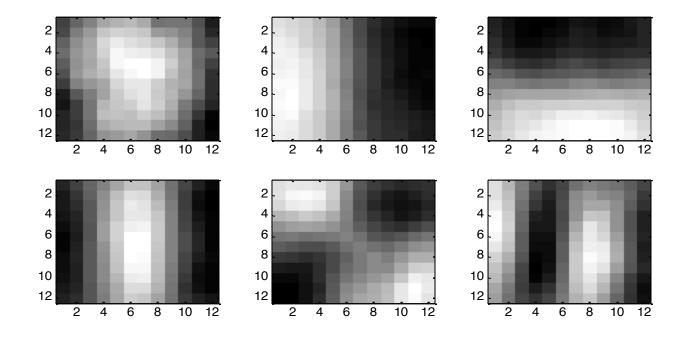
16 most important eigenvectors



PCA compression: 144D) 6D



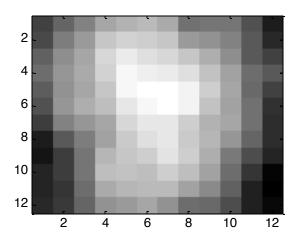
6 most important eigenvectors

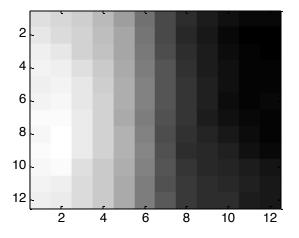


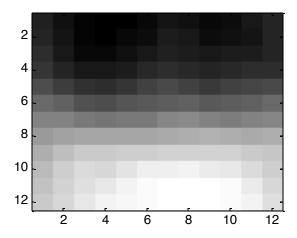
PCA compression: 144D) 3D



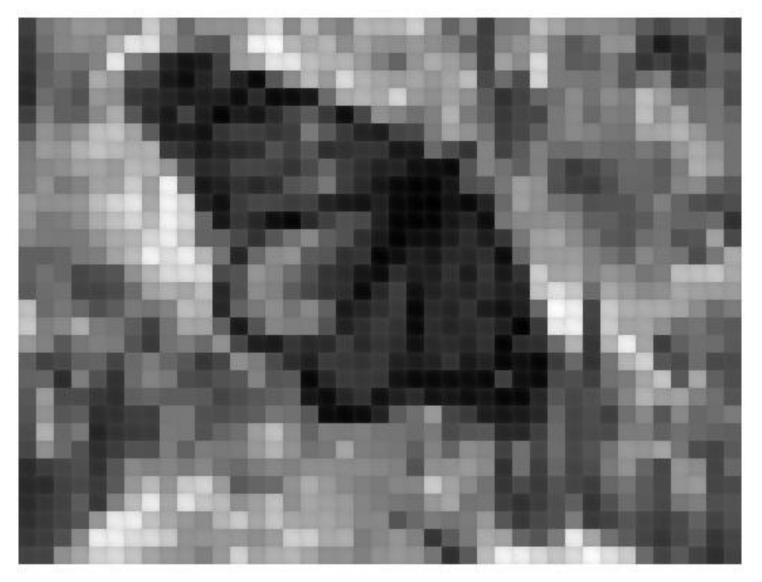
3 most important eigenvectors



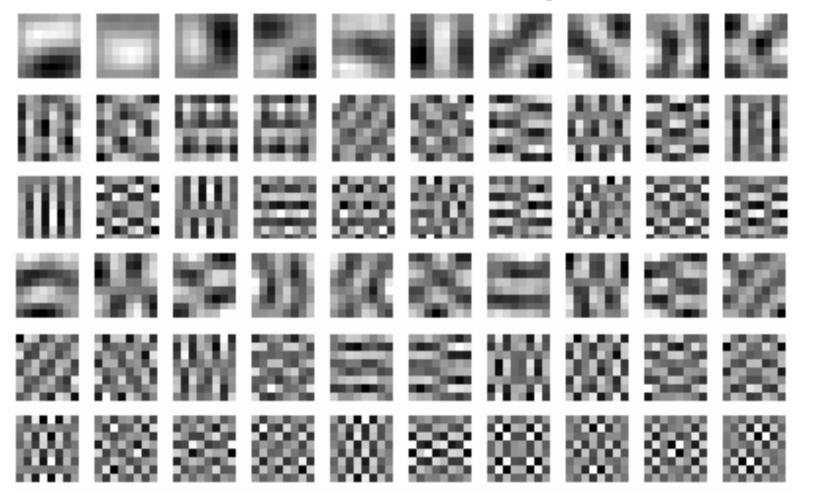




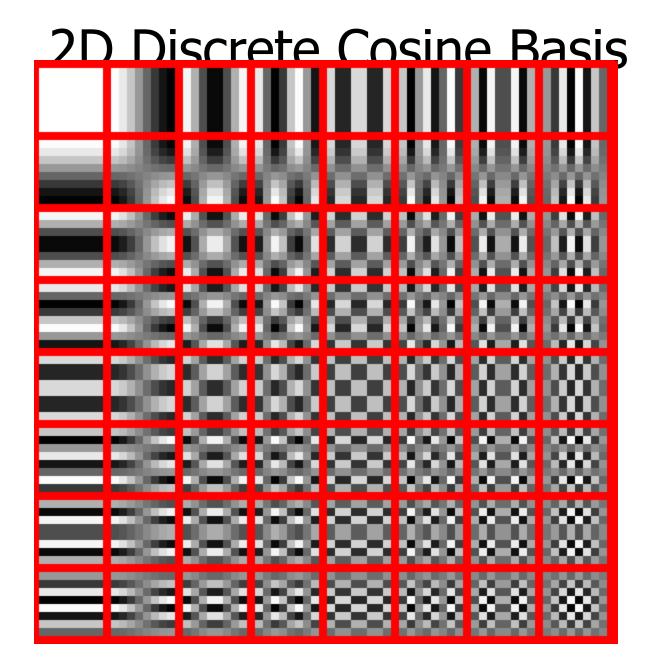
PCA compression: 144D) 1D



60 most important eigenvectors



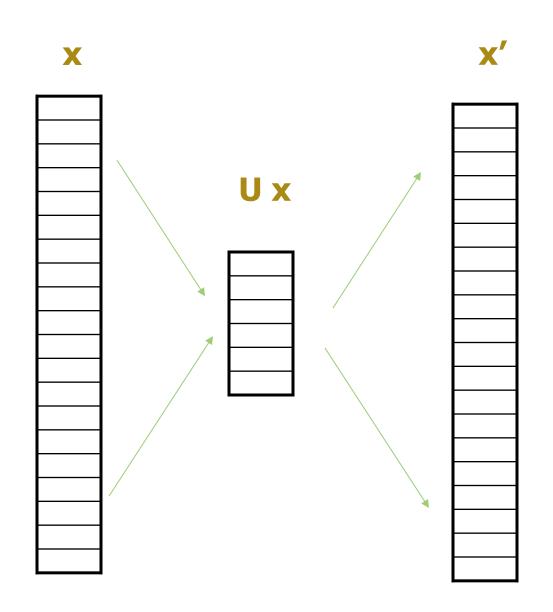
Looks like the discrete cosine bases of JPG!...



http://en.wikipedia.org/wiki/Discrete_cosine_transform

Noise Filtering

Noise Filtering, Auto-Encoder...



Noisv image



Denoised image using 15 PCA components

