Piyush Rai

CS5350/6350: Machine Learning

October 20, 2011

(CS5350/6350)

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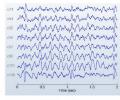
High-Dimensional Datasets Abound ...



face images

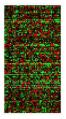
Zambian President Levy Mwanawasa has won a second term in office in an election his challenger Michael Sata accused him of rigging, official results showed on Monday. According to media reports, a pair of hackens said on Saturday that the Firefox Web browser, commonly perceived as the safer and more customizable alternative to market leader Internet Explorer, is critically flawed. A presentation on the flaw was shown during the ToorCon hacker conference in San Diego.

documents



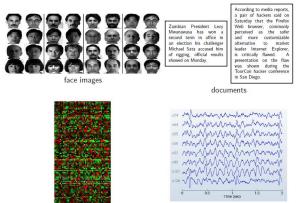
MEG readings

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gene expression data

High-Dimensional Datasets Abound ...



MEG readings

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Goal: Find a low-dimensional, yet useful representation of the data

gene expression data

(CS5350/6350)

Linear Dimensionality Reduction

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• Insights into the low-dimensional structures in the data (visualization)

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- Fewer dimensions \Rightarrow Less chances of overfitting \Rightarrow Better generalization

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- Note: Dimensionality Reduction is different from Feature Selection
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- Fewer dimensions \Rightarrow Less chances of overfitting \Rightarrow Better generalization
- Speeding up learning algorithms
 - Most algorithms scale badly with increasing data dimensionality
- Less storage requirements (data compression)
- Note: Dimensionality Reduction is different from Feature Selection
 ... although the goals are kind of the same
- Dimensionality reduction is more like "Feature Extraction"
 - Constructing a small set of new features from the original features

• Based on the idea of doing a linear projection of the data

- Based on the idea of doing a linear projection of the data
- Works well if the data lies close to a linear subspace

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- Based on the idea of doing a linear projection of the data
- Works well if the data lies close to a linear subspace
- Consider a high dimensional example $\mathbf{x} \in \mathbb{R}^D$
- We want to project it down to a K-dimensional vector \mathbf{z} (K \ll D) $\mathbf{z} = \mathbf{U}^{\top} \mathbf{x}$
- $\mathbf{z} \in \mathbb{R}^{K}$ is the projection
- **U** is the $D \times K$ projection matrix (defining K projection directions)

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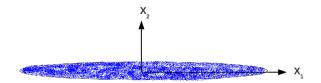
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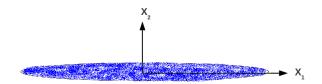


- $\bullet\,$ Different methods differ in how ${\bf U}$ is defined/learned
- The differences depend on what properties of data we want to capture

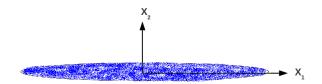


- Consider this 2 dimensional data
- Each example **x** has 2 features $\{x_1, x_2\}$

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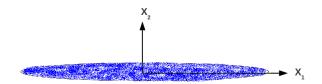


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- Consider ignoring the feature x₂ for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$



- Consider this 2 dimensional data
- Each example **x** has 2 features {*x*₁, *x*₂}
- Consider ignoring the feature x₂ for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$
- Are we losing much information by throwing away x₂?

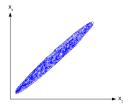
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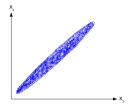
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- Each example **x** has 2 features $\{x_1, x_2\}$
- Consider ignoring the feature x₂ for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$
- Are we losing much information by throwing away x_2 ?
- No. Most of the data spread is along x_1 (very little variance along x_2)

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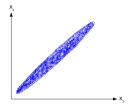
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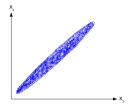


- Consider this 2 dimensional data
- Each example **x** has 2 features $\{x_1, x_2\}$
- Consider ignoring the feature x₂ for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$



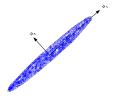
- Consider this 2 dimensional data
- Each example **x** has 2 features $\{x_1, x_2\}$
- Consider ignoring the feature x_2 for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$
- Are we losing much information by throwing away x₂?

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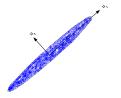
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- Consider ignoring the feature x₂ for each example
- Each 2-dimensional example **x** now becomes 1-dimensional $\mathbf{x} = \{x_1\}$
- Are we losing much information by throwing away x_2 ?
- Yes. The data has substantial variance along both features (i.e., both axes)

A D > A P > A B > A



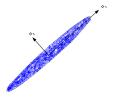
• Now consider a change of axes (the co-ordinate system)

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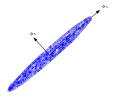
- Now consider a change of axes (the co-ordinate system)
- Each example **x** has 2 features $\{u_1, u_2\}$

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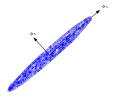


- Now consider a change of axes (the co-ordinate system)
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- No. Most of the data spread is along u_1 (very little variance along u_2)

(CS5350/6350)

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- PCA: Take top K PC's and project the data along those

(a)

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 - Orthogonal: $\mathbf{u}_i^\top \mathbf{u}_j = 0$ if $i \neq j$, Orthonormal: $\mathbf{u}_i^\top \mathbf{u}_i = 1$

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- Each principal component is a vector of size $D \times 1$
- We want only the first K principal components

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where \boldsymbol{S} is the data covariance matrix defined as

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- Want to have **u**₁ that maximizes the projected data variance **u**₁⁺**Su**₁
 - Subject to the constraint: $\mathbf{u}_1^\top \mathbf{u}_1 = 1$
 - We will introduce a Lagrange multiplier λ_1 for this constraint

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- Objective function: $\mathbf{u}_1^{\top} \mathbf{S} \mathbf{u}_1 + \lambda_1 (1 \mathbf{u}_1^{\top} \mathbf{u}_1)$
- Taking derivative w.r.t. **u**₁ and setting it to zero gives:

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- We know that the projected data variance $\mathbf{u}_1^\top \mathbf{S} \mathbf{u}_1 = \lambda_1$ is maximum
 - Thus λ_1 should be the largest eigenvalue
 - Thus \mathbf{u}_1 is the first (top) eigenvector of **S** (with eigenvalue λ_1)
 - \Rightarrow the first principal component (direction of highest variance in the data)

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 - Thus u₁ is the first (top) eigenvector of S (with eigenvalue λ₁)
 ⇒ the first principal component (direction of highest variance in the data)
- Subsequent PC's are given by the subsequent eigenvectors of ${\boldsymbol{\mathsf{S}}}$

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PCA: The Algorithm

• Compute the mean of the data

$$\bar{\mathbf{x}} = \frac{1}{N} \sum_{n=1}^{N} \mathbf{x}_n$$

• Compute the sample covariance matrix (using the mean subtracted data)

$$\mathbf{S} = \frac{1}{N} \sum_{n=1}^{N} (\mathbf{x}_n - \bar{\mathbf{x}}) (\mathbf{x}_n - \bar{\mathbf{x}})^{\top}$$

- Do the eigenvalue decomposition of the $D \times D$ matrix **S**
- Take the top K eigenvectors (corresponding to the top K eigenvalues)
- Call these $\mathbf{u}_1, \ldots, \mathbf{u}_K$ (s.t. $\lambda_1 \ge \lambda_2 \ge \ldots \lambda_{K-1} \ge \lambda_K$)
- $\mathbf{U} = [\mathbf{u}_1 \ \mathbf{u}_2 \ \dots \ \mathbf{u}_K]$ is the projection matrix of size $D \times K$

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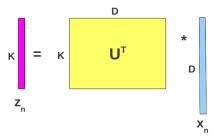
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- Projection of each example \mathbf{x}_n is computed as $\mathbf{z}_n = \mathbf{U}^\top \mathbf{x}_n$
 - \mathbf{z}_n is a $K \times 1$ vector (also called the embedding of \mathbf{x}_n)

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PCA: Pictorially

• For a single example \mathbf{x}_n :

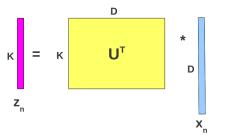


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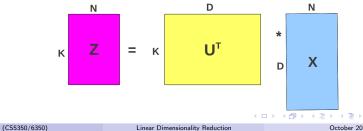
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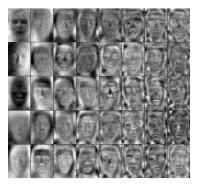
• For a set of *N* examples:



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PCA Example: Eigenfaces

• Principal Components learned using a face image dataset



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PCA: Approximate Reconstruction

Given the principal components u₁,..., u_K, the PCA approximation of an example x_n is:

$$\tilde{\mathbf{x}}_n = \sum_{i=1}^K (\mathbf{x}_n^\top \mathbf{u}_i) \mathbf{u}_i = \sum_{i=1}^K z_{ni} \mathbf{u}_i$$

where $\mathbf{z}_n = [z_{n1}, \dots, z_{nK}]$ is the low-dimensional projection of \mathbf{x}_n

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- This gives us a way of compressing data
- To compress a dataset $\mathbf{X} = [\mathbf{x}_1, \dots, \mathbf{x}_N]$, all we need is the set of $K \ll D$ principal components, and the projections $\mathbf{Z} = [\mathbf{z}_1, \dots, \mathbf{z}_N]$ of each example

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- Recall: PCA requires eigen-decomposition of $D \times D$ covariance matrix $\mathbf{S} = \frac{1}{N} \mathbf{X} \mathbf{X}^{\top}$ (assuming centered data, and \mathbf{X} being $D \times N$)
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- The eigenvectors aren't exactly the same (but still related)
- The relationship is $\mathbf{u}_i = \frac{1}{(N\lambda_i)^2} \mathbf{X} \mathbf{v}_i$
- { λ_i , \mathbf{v}_i } is an eigenvalue-eigenvector pair of the $N \times N$ matrix $\frac{1}{N} \mathbf{X}^{\top} \mathbf{X}$, and \mathbf{u}_i is the corresponding eigenvector of $\mathbf{S} = \frac{1}{N} \mathbf{X} \mathbf{X}^{\top}$ (that we want)

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• Dimensionality reduction with label information (when the ultimate goal is classification/regression)

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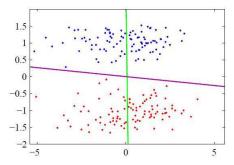
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• PCA: magenta line, FDA: green line



PCA based projection makes the classes overlap (which is bad)
LDA/FDA is often better if the final goal is classification

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