# Linear Dimensionality Reduction 

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CS5350/6350: Machine Learning
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## High-Dimensional Datasets Abound..


documents


MEG readings
gene expression data

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

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- 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


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- Consider a high dimensional example $\mathbf{x} \in \mathbb{R}^{D}$
- We want to project it down to a $K$-dimensional vector $\mathbf{z}(K \ll D)$

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\mathbf{z}=\mathbf{U}^{\top} \mathbf{x}
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- $\mathbf{z} \in \mathbb{R}^{K}$ is the projection
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- Different methods differ in how $\mathbf{U}$ is defined/learned
- The differences depend on what properties of data we want to capture


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- Yes. The data has substantial variance along both features (i.e., both axes)


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- Note: All principal components are orthogonal to each other
- PCA: Take top K PC's and project the data along those


## PCA: Finding the Principal Components

- Given: $N$ examples $\mathbf{x}_{1}, \ldots, \mathbf{x}_{N}$, each example $\mathbf{x}_{n} \in \mathbb{R}^{D}$
- Goal: Project the data from $D$ dimensions to $K$ dimensions $(K<D)$
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- Want to capture the maximum possible variance in the projected data
- Let $\mathbf{u}_{1}, \ldots, \mathbf{u}_{D}$ be the principal components, assumed to be:
- 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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- We want only the first $K$ principal components


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\frac{1}{N} \sum_{n=1}^{N}\left\{\mathbf{u}_{1}^{\top} \mathbf{x}_{n}-\mathbf{u}_{1}^{\top} \overline{\mathbf{x}}\right\}^{2}=\mathbf{u}_{1}^{\top} \mathbf{S} \mathbf{u}_{1}
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where $\mathbf{S}$ is the data covariance matrix defined as

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- Want to have $\mathbf{u}_{1}$ that maximizes the projected data variance $\mathbf{u}_{1}^{\top} \mathbf{S} \mathbf{u}_{1}$
- Subject to the constraint: $\mathbf{u}_{1}^{\top} \mathbf{u}_{1}=1$
- We will introduce a Lagrange multiplier $\lambda_{1}$ for this constraint


## PCA: Finding the Principal Components

- Objective function: $\mathbf{u}_{1}^{\top} \mathbf{S} \mathbf{u}_{1}+\lambda_{1}\left(1-\mathbf{u}_{1}^{\top} \mathbf{u}_{1}\right)$
- Taking derivative w.r.t. $\mathbf{u}_{1}$ 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 $\lambda_{1}$ should be the largest eigenvalue
- Thus $\mathbf{u}_{1}$ is the first (top) eigenvector of $\mathbf{S}$ (with eigenvalue $\lambda_{1}$ ) $\Rightarrow$ the first principal component (direction of highest variance in the data)


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- Subsequent PC's are given by the subsequent eigenvectors of $\mathbf{S}$


## PCA: The Algorithm

- Compute the mean of the data

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\overline{\mathrm{x}}=\frac{1}{N} \sum_{n=1}^{N} \mathrm{x}_{n}
$$

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

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\mathbf{S}=\frac{1}{N} \sum_{n=1}^{N}\left(\mathbf{x}_{n}-\overline{\mathbf{x}}\right)\left(\mathbf{x}_{n}-\overline{\mathbf{x}}\right)^{\top}
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- Do the eigenvalue decomposition of the $D \times D$ matrix $\mathbf{S}$
- Take the top $K$ eigenvectors (corresponding to the top $K$ eigenvalues)
- Call these $\mathbf{u}_{1}, \ldots, \mathbf{u}_{K}$ (s.t. $\lambda_{1} \geq \lambda_{2} \geq \ldots \lambda_{K-1} \geq \lambda_{K}$ )
- $\mathbf{U}=\left[\begin{array}{llll}\mathbf{u}_{1} & \mathbf{u}_{2} & \ldots & \mathbf{u}_{K}\end{array}\right]$ is the projection matrix of size $D \times K$


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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 $\mathbf{x}_{n}$ )


## PCA: Pictorially

- For a single example $\mathbf{x}_{n}$ :



## PCA: Pictorially

- For a single example $\mathbf{x}_{n}$ :

- For a set of $N$ examples:



## PCA Example: Eigenfaces

- Principal Components learned using a face image dataset



## PCA: Approximate Reconstruction

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## X (DxN) <br> U (DxK)

$($ ? $\ldots$ 国 $) \approx($ ?

## PCA for Very High Dimensional Data

- In many cases, $N<D$
- 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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- Fact: $\mathbf{S}=\frac{1}{N} \mathbf{X} \mathbf{X}^{\top}$ has the same $N-1$ non-zero eigenvalues as that of the $N \times N$ matrix $\frac{1}{N} \mathbf{X}^{\top} \mathbf{X}$ (for which eigen-decomposition is cheaper if $N<D$ )


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- The remaining $D-N+1$ eigenvalues are zero
- Fact: $\mathbf{S}=\frac{1}{N} \mathbf{X} \mathbf{X}^{\top}$ has the same $N-1$ non-zero eigenvalues as that of the $N \times N$ matrix $\frac{1}{N} \mathbf{X}^{\top} \mathbf{X}$ (for which eigen-decomposition is cheaper if $N<D$ )
- The eigenvectors aren't exactly the same (but still related)
- The relationship is $\mathbf{u}_{i}=\frac{1}{\left(N \lambda_{i}\right)^{2}} \mathbf{X} \mathbf{v}_{i}$
- $\left\{\lambda_{i}, \mathbf{v}_{i}\right\}$ 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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- PCA ignores label information even if it is available
- Only chooses directions of maximum variance
- Fisher Discriminant Analysis (FDA) takes into account the label information
- It's also called Linear Discriminant Analysis (LDA)


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## PCA vs FDA/LDA

- 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

