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# Lecture 12: Face Recognition & Dimensionality Reduction

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## 1 Overview and Motivation

### 1.1 Overview

Dimensionality reduction is a process for reducing the number of features to be used in an analysis or model. There are several methods commonly used in dimensionality reduction. The two main methods covered this lecture are Singular Value Decomposition (SVD) and Principal Component Analysis (PCA).

### 1.2 Motivation

There are several reasons why reducing the dimensions for a model is useful.

1. Reduction in computational cost can be achieved. In many data sets, most of the variance can be explained by a relatively small number of principal components, when compared to the dimensionality of the entire feature set. By reducing our dimensions to focus on these key components, we can shrink the computational cost of analyzing each data point without losing much granularity in the data.
2. Reduce the effects of the “curse of dimensionality”. In lecture 11 we learned that as we increase the dimension of a feature space, the number of data points needed to “fill in” that space with the same density explodes exponentially. That is to say, the more dimensions there are, the more examples a machine learning algorithm needs to learn from and the longer the algorithm must run to analyze the same number of data points. By performing dimensionality reduction, we can mitigate the effects of this “curse of dimensionality”.
3. Compress data. By shrinking the dimensionality of an image, we can dramatically reduce the amount of storage required to store the image.  
In such cases the computational cost per data point may be reduced by many orders of magnitude with a procedure like SVD

## 2 Singular Value Decomposition

### 2.1 Overview

Intuitively, Singular Value Decomposition (SVD) is a procedure that allows one to (in a sense) “rotate the axes” of the feature space in which a dataset has been plotted so that instead of the different axes corresponding to specific features (e.g. age, income, gender, etc. of a customer) the new axes point in directions that track linear combinations of the original feature space. These “new axes” are useful

because they systematically break down the variance in the data points (how widely the data points are distributed) based on each directions contribution to the variance in the data:

The result of this process is a ranked list of "directions" in the feature space ordered from most variance to least. The directions along which there is greatest variance are referred to as the "principal components" (of variation in the data) and the common wisdom is that by focusing on the way the data is distributed along these dimensions exclusively, one can capture most of the information represented in the original feature space without having to deal with such a high number of dimensions (but see below on the difference between feature selection and dimensionality reduction).

## 2.2 Technical Details of Singular Value Decomposition

- SVD represents any matrix A as a product of three matrices:  $A = U\Sigma V^T$  where U and V are rotation matrices, and  $\Sigma$  is a diagonal scaling matrix. For example:

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

- For many readers, it may be sufficient to extract SVD values by writing: `[U, S, V] = numpy.linalg.svd(A)`. However, the underpinnings of how SVD is computed is useful for later topics. Computers typically compute SVD by taking the following steps:
  - Compute the eigenvectors of  $AA^T$ . These vectors are the columns of U. Square root of the eigenvalues are the singular values (entries of  $\Sigma$ )
  - Compute the eigenvectors of  $A^T A$ . These vectors are columns of V (or rows of  $V^T$ )
- Since SVD relies on eigenvector computation, which are typically fast, SVD can be performed quite quickly; even for large matrices.
- A more detailed, geometric explanation of SVD may be found [here](#)[1].

## 2.3 Applications of Singular Value Decomposition

- One the most utilized applications of SVD is the computation of matrix inverses. If an arbitrary matrix A can be decomposed by way of:  $A = U\Sigma V^T$ , the inverse of A may be defined as:  $A^+ = V^T \Sigma^{-1} U$ . Although this inverse is an approximation, it allows one to calculate the inverses of many non-square matrices. MacAusland (2014) discusses the mathematical basis of this inverse, which is named the Moore-Penrose inverse, after its creators[3]. Unsurprisingly, a large variety of matrix problems can be solved by utilizing this approach.
- Singular Value Decomposition can also be used to compute the Principal Components of a matrix. Principal Components are heavily utilized in various data analysis and machine learning routines, hence SVD is typically a core routine within many programs.

# 3 Principal Component Analysis

## 3.1 What are Principal Components?

Continuing with the SVD example we have above, notice that Column 1 of U gets scaled by the first value from  $\Sigma$ .

$$\begin{matrix} U\Sigma & & V^T & & A_{\text{partial}} \\ \begin{bmatrix} -3.67 & -.71 & 0 \\ -8.8 & .30 & 0 \end{bmatrix} & \times & \begin{bmatrix} -.42 & -.57 & -.70 \\ .81 & .11 & -.58 \\ .41 & -.82 & .41 \end{bmatrix} & & \begin{bmatrix} 1.6 & 2.1 & 2.6 \\ 3.8 & 5.0 & 6.2 \end{bmatrix} \end{matrix}$$

Then, the resulting vector  $U\Sigma$  gets scaled by row 1 of  $V^T$  to produce a contribution to the columns of A which is denoted  $A_{\text{partial}}$ . Each product of (column i of U)\*(value i from  $\Sigma$ )\*(row i of  $V^T$ )

produces a component of the final A.

$$\begin{aligned}
 & \begin{matrix} U\Sigma \\ \begin{bmatrix} -3.67 & -.71 & 0 \\ -8.8 & .30 & 0 \end{bmatrix} \end{matrix} \times \begin{matrix} V^T \\ \begin{bmatrix} -.42 & -.57 & -.70 \\ .81 & .11 & -.58 \\ .41 & -.82 & .41 \end{bmatrix} \end{matrix} = \begin{matrix} A_{\text{partial}} \\ \begin{bmatrix} 1.6 & 2.1 & 2.6 \\ 3.8 & 5.0 & 6.2 \end{bmatrix} \end{matrix} \\
 + & \begin{matrix} U\Sigma \\ \begin{bmatrix} -3.67 & -.71 & 0 \\ -8.8 & .30 & 0 \end{bmatrix} \end{matrix} \times \begin{matrix} V^T \\ \begin{bmatrix} -.42 & -.57 & -.70 \\ .81 & .11 & -.58 \\ .41 & -.82 & .41 \end{bmatrix} \end{matrix} = \begin{matrix} A_{\text{partial}} \\ \begin{bmatrix} -.6 & -.1 & .4 \\ .2 & 0 & -.2 \end{bmatrix} \end{matrix} \\
 = & \begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix}
 \end{aligned}$$

In this process we are building the the matrix A as a linear combination of the columns of U. As seen above, if we use all columns of U, we rebuild A perfectly. However, in real-world data, **we can use only the first few columns of U** to get a good approximation of A. This arises due to the properties that  $\Sigma$  has.  $\Sigma$  is a diagonal matrix where the largest value is in the top left corner, and the rest of the values on the diagonal decreases as you move to the right. Thus, the first few columns of  $U$  contribute the largest weight towards  $A$ . These first few columns of  $U$  are called **principal components**.

However, not all matrices can be easily compressed as in the previous example. One way to figure out if it's worth performing is through Principal Component Analysis. From a high level standpoint, we want to see if it's possible to remove dimensions that don't contribute much to the final image. We do this by analyzing the covariance matrix. Although the value of covariance doesn't matter as much, the sign of covariance does, with positive indicating positive correlation and negative indicating negative correlation. A covariance of zero indicates the two are independent of one another.

### 3.2 Performing Principal Component Analysis

Principal Component Analysis can be performed using the sklearn package: `sklearn.decomposition.PCA`. However, it was alluded to earlier that SVD can be used to perform Principal Component Analysis. A non-formal approach is outlined below:

1. Format your data into a  $m * n$  matrix where  $m$  denotes the number of samples and  $p$  represents the number of features or variables corresponding to a single sample.
2. Center the matrix  $X$  by subtracting the mean and dividing by the standard deviation along each column(feature) in  $X$
3. Diagonalizing  $X$  using SVD yields:  $X = U\Sigma V^T$
4. Eigenvectors are the principal directions and the projections on these axes are the components. This ultimately means we want to compute  $XV$
5. Since  $V$  holds eigenvectors and is thus orthonormal,  $XV = U\Sigma V^T V = US$
6. (5) implies we simply need the columns of  $US$ , both matrices that are surfaced by SVD

Detailed explanations that elucidate the reasoning behind the above steps are discussed by Moore (1981) and can be found on numerous websites online.

### 3.3 Applications of Principal Components

- PCA has been extensively used in image compression. Much of the information captured within an image matrix can be extracted using matrices of lower ranks. This allows large images to be compressed without significant loss of quality. An example of PCA based compression, using only the first 16 principal components, is shown below:



Figure 1: Right: Original Image, Left: Compressed Image

With just the first 16 principal components, an image that closely resembles the original image can be reconstructed. The relative error as a result of the dimensions used for PCA for the above image is shown below:

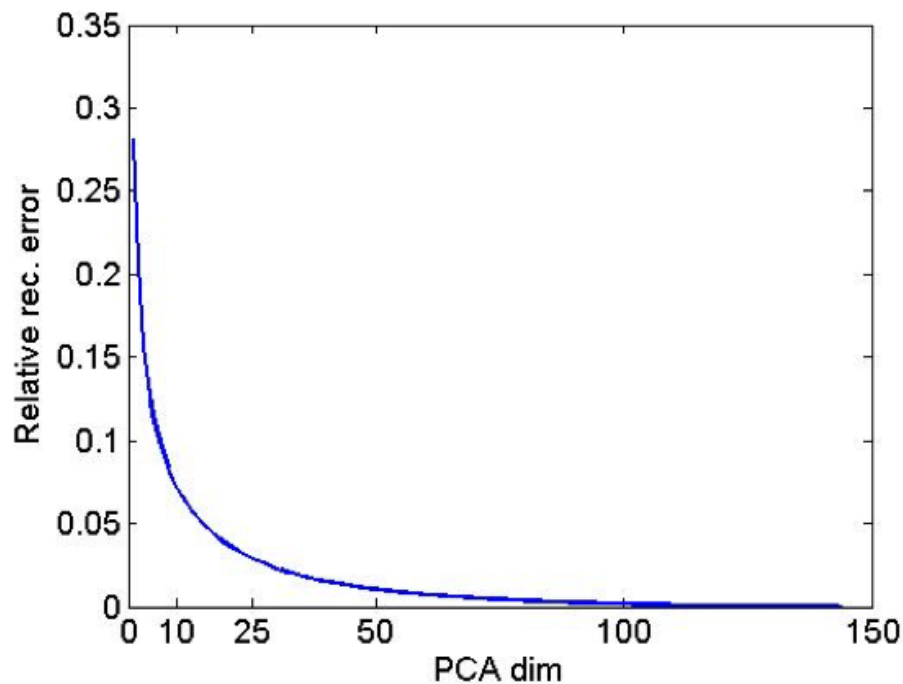


Figure 2: Relative Error as Function of PCA dimensions

- Web search engines also utilize PCA. There are billions of pages on the Internet that may have a non-trivial relation to a provided search phrase. Companies such as Google, Bing and Yahoo typically narrow the search space by only considering a small subset of this search matrix, which can be extracted using PCA[2]. This is critical for timely, efficient searches and speaks to the power of SVD.

In essence, PCA allows you to represent data samples as weights on various components - allowing one to essentially represent the difference between samples. This can significantly reduce data redundancy and can make algorithms used in a variety of industries more efficient and insightful!

## References

- [1] <http://www.ams.org/samplings/feature-column/fcarc-svd>.
- [2] Snasel V. Abdulla H.D. Search result clustering using a singular value decomposition (svd). 2009.
- [3] R. MacAusland. The moore-penrose inverse and least squares. 2014.