

Dimensional Reduction

Principal Component Analysis



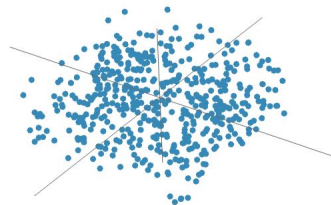


Goals

- Understand the need and purpose of dimensionality reduction algorithms.
- Understand and learn the details of Principal Component Analysis (PCA). Including its strengths and weaknesses.
- See concrete applications of using PCA in context.
- Show other dimensionality reduction algorithms.



Point Cloud Data



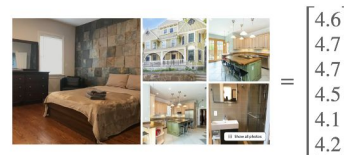
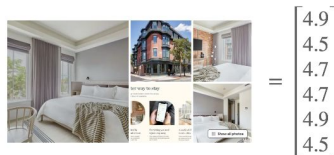
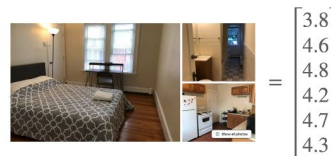
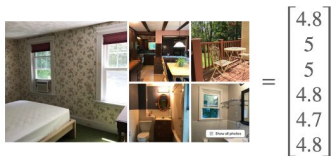
A point cloud is a collection of data points in \mathbb{R}^n

- Each point is represented according to its coordinates $(X_1, X_2, X_3, \dots, X_m)$
- Each coordinate may represent a different feature or characteristic:
 - Hotels in a city can be represented in \mathbb{R}^6 according to user rating of characteristics: cleanliness, accuracy, communication, check-in, location, value
 - A 28x28 pixel grayscale image can be represented in \mathbb{R}^{784} : each pixel is represented with a unique number according to a scale from black to white
 - Samples of expressions of N genes can be represented in \mathbb{R}^N

Example 1: Hotel Listings

- Each data point in \mathbb{R}^6 corresponds to a hotel
- Hotels are ranked according to 6 categories:
- Each individual hotel can be represented by a point in

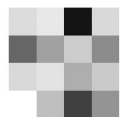
Cleanliness	_____ 4.8	Accuracy	_____ 4.8
Communication	_____ 5.0	Location	_____ 4.7
Check-in	_____ 5.0	Value	_____ 4.8





Example 2: Grayscale Images

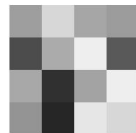
- Each data point corresponds to an image of resolution 4x4
- Each of the 16 pixels is represented with a number from 0 (black) to 1 (white)
- Each image can be represented by a point in \mathbb{R}^{16}



$$\begin{pmatrix} 0.86 & 0.91 & 0.08 & 0.85 \\ 0.4 & 0.63 & 0.8 & 0.55 \\ 0.85 & 0.89 & 0.68 & 0.79 \\ 1. & 0.75 & 0.25 & 0.58 \end{pmatrix}$$



$$\begin{pmatrix} 0.86 \\ 0.91 \\ 0.08 \\ 0.85 \\ 0.4 \\ 0.63 \\ 0.8 \\ 0.55 \\ 0.85 \\ 0.89 \\ 0.68 \\ 0.79 \\ 1. \\ 0.75 \\ 0.25 \\ 0.58 \end{pmatrix}$$



$$\begin{pmatrix} 0.62 & 0.84 & 0.65 & 0.61 \\ 0.3 & 0.67 & 0.93 & 0.35 \\ 0.66 & 0.19 & 0.64 & 0.93 \\ 0.58 & 0.15 & 0.9 & 0.84 \end{pmatrix}$$



$$\begin{pmatrix} 0.62 \\ 0.84 \\ 0.65 \\ 0.61 \\ 0.3 \\ 0.67 \\ 0.93 \\ 0.35 \\ 0.66 \\ 0.19 \\ 0.64 \\ 0.93 \\ 0.58 \\ 0.15 \\ 0.9 \\ 0.84 \end{pmatrix}$$



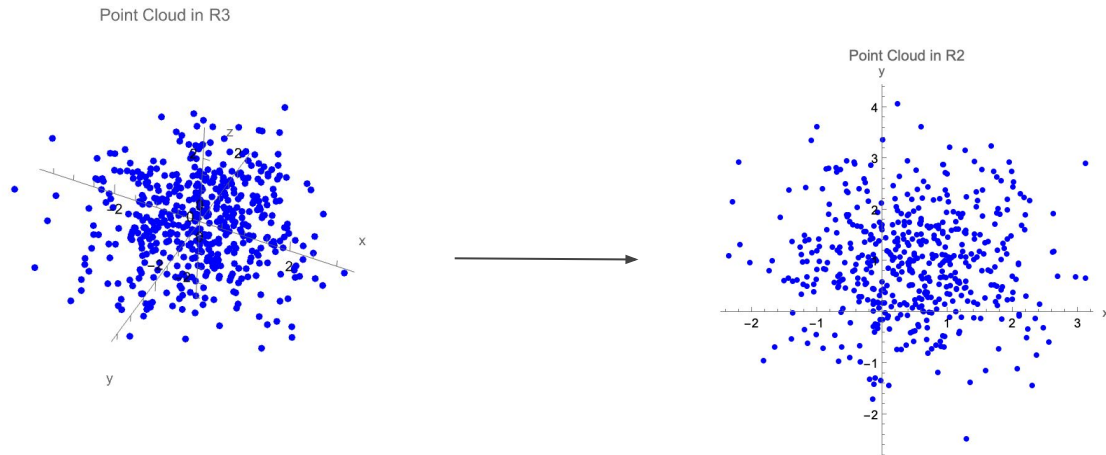
Example 3: Gene Expression

- Each sample measures expressions of N genes in distinct cells
- Each individual cell can be represented by a point in \mathbb{R}^N

Sample ID	Gene1	Gene2	Gene3	...	GeneN
Sample1	5.2	0.1	3.4	...	7.6
Sample2	4.9	0.0	3.8	...	6.8

Point Cloud Data: Goal

- Is there a way to visualize higher dimensional data?
- If so, how much is it representative of the original data?
- Are there any features that are more important than others?
- Are there any **combinations of features** that are more important than others?





Why Dimensional Reduction?

Reasons are practical from both a computational and statistical point of view:

- Reduce computational complexity
- Reduce redundancy and noise
- Reduce overfitting (improve performance)
- Enable visualization
- Find correlations between input features

What do we need?

- Statistics
- Linear Algebra



Statistics Basics



Statistics Measurements

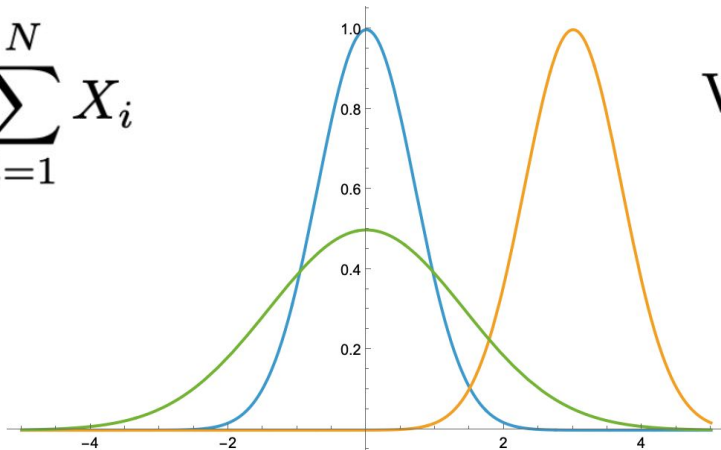
Suppose we have N measurements of a certain feature: X_1, X_2, \dots, X_N

The **mean** is the central tendency or "average" of a set of numbers:

The **variance** measures how spread out the values are around the mean:

$$\bar{X} = \frac{1}{N} \sum_{i=1}^N X_i$$

$$\text{Var}(X) = \frac{1}{N} \sum_{i=1}^N (X_i - \bar{X})^2$$





Statistics Measurements

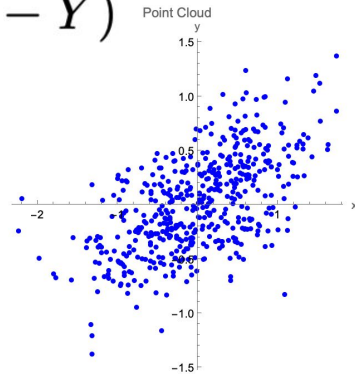
Suppose we have N measurements of two features:

$$X_1, X_2, \dots, X_N$$
$$Y_1, Y_2, \dots, Y_N$$

The **covariance** is a measure of how two variables change together—whether they tend to increase or decrease at the same time.

$$\text{Cov}(X, Y) = \frac{1}{N} \sum_{i=1}^N (X_i - \bar{X})(Y_i - \bar{Y})$$

Correlation is the standardized version of covariance





Linear Algebra Basics



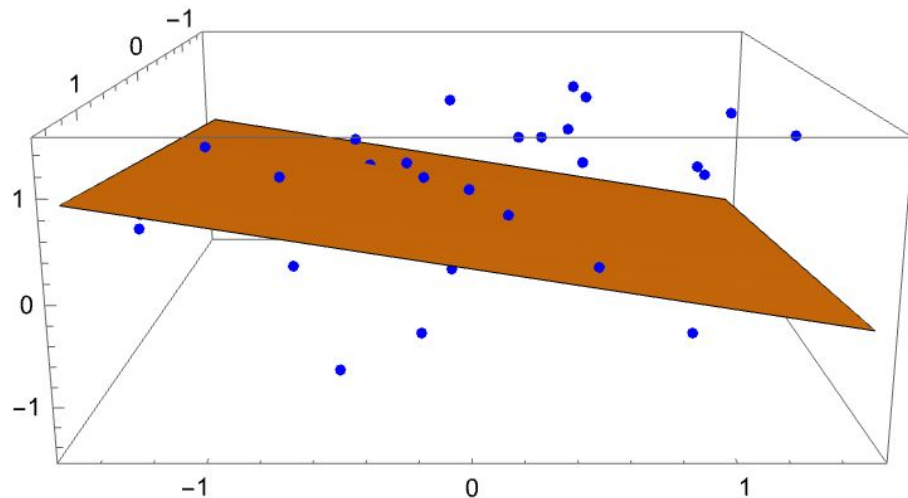
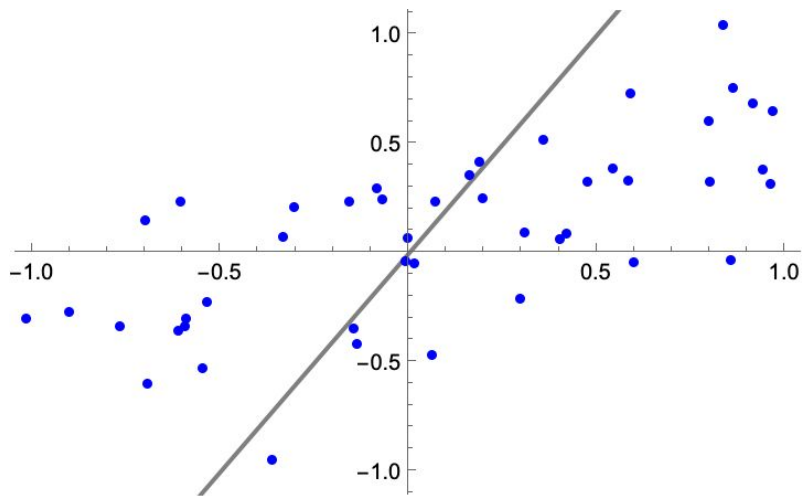
Linear Transformations

Suppose we have N measurements of a certain feature: X_1, X_2, \dots, X_N



Orthogonal Projections

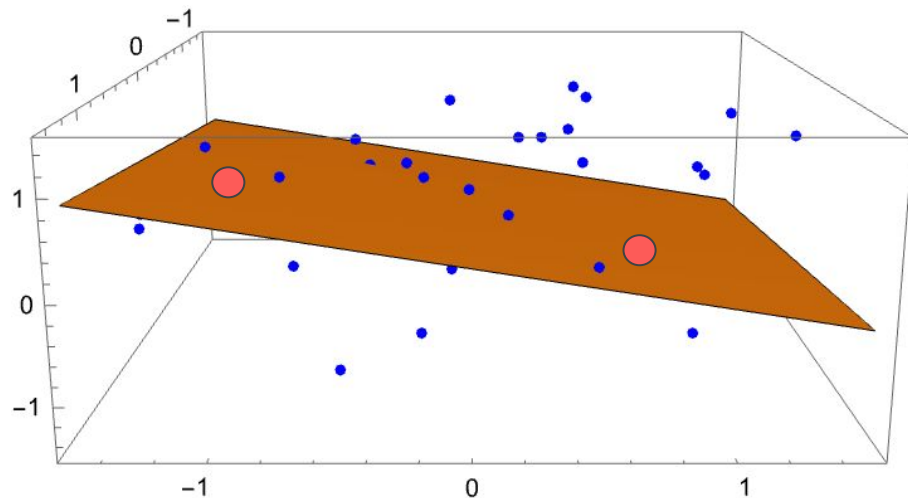
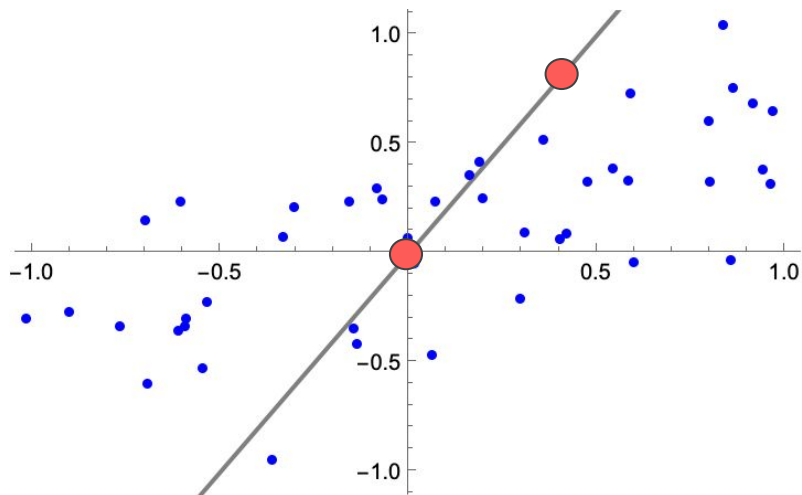
- Orthogonal Projections are a type of Linear Transformation
- Let V be a n -dimensional subspace of \mathbb{R}^n (line in \mathbb{R}^2 , line or plane in \mathbb{R}^3 , etc.)
- Orthogonal projections minimize distance between points and projections.





Orthogonal Projections

- Orthogonal Projections have special points:
 - Points who's value doesn't chage
 - Points who's value becomes 0





Eigenvalues & Eigenvectors

Eigenvectors and eigenvalues are specific properties of square ($n \times n$) matrices.
Definition is not super important.

Suppose A is an $x \times x$ matrix. A nonzero vector \vec{v} in \mathbb{R}^n is an **eigenvector** of A of **eigenvalue** λ if

$$A\vec{v} = \lambda\vec{v}.$$

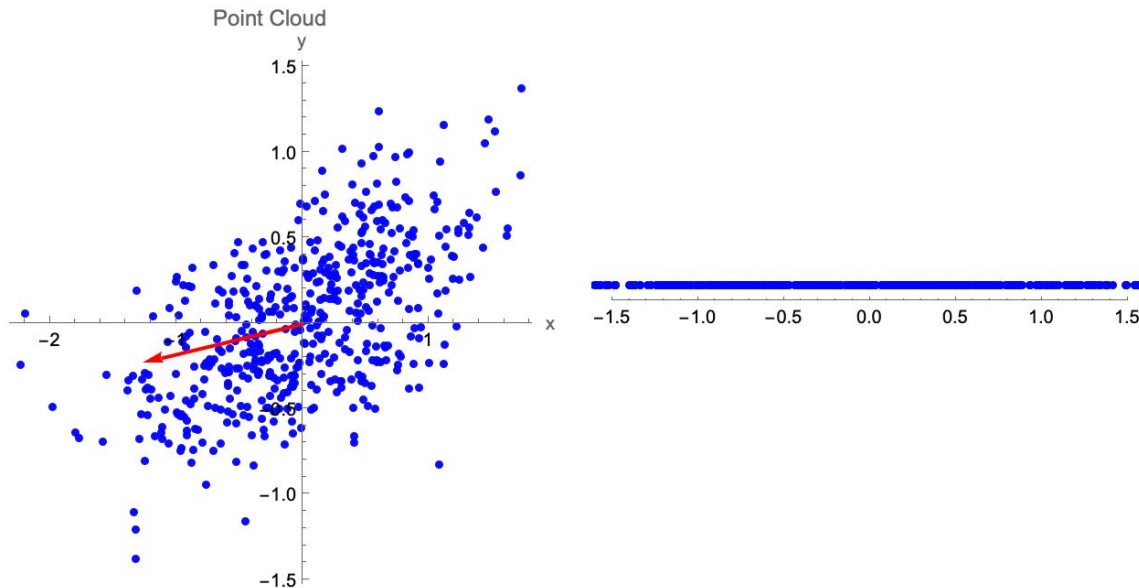


Principal Component Analysis



Principal Component Analysis

- PCA finds the direction(s) in which the data varies the most (i.e., is most spread out), and
- projects the data onto those directions to reduce dimensionality while preserving as much variance as possible.





Step 0: Gathering The Data

- Consider a multidimensional dataset consisting of **N** observations of **m** different characteristics X_i :

$$(X_1^{(1)}, X_2^{(1)}, X_3^{(1)}, \dots, X_m^{(1)})$$

$$(X_1^{(2)}, X_2^{(2)}, X_3^{(2)}, \dots, X_m^{(2)})$$

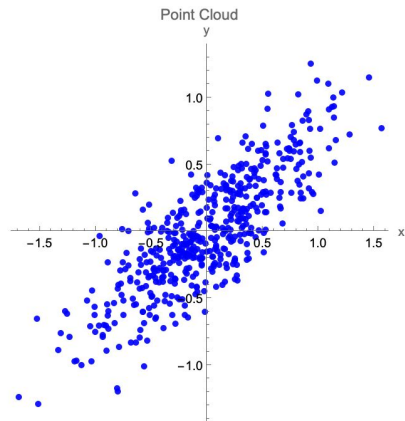
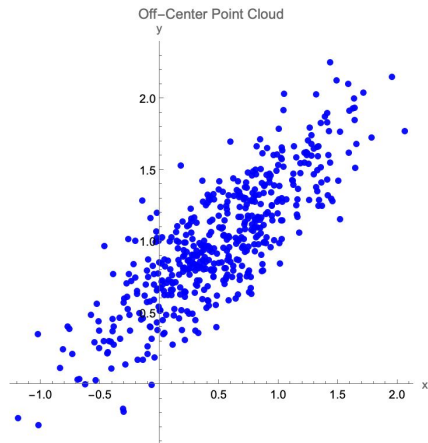
$$\vdots$$

$$(X_1^{(N)}, X_2^{(N)}, X_3^{(N)}, \dots, X_m^{(N)})$$

- This data lives in a high-dimensional space \mathbb{R}^m that is “impossible” for us to visualize

Step 1: Standardizing The Data

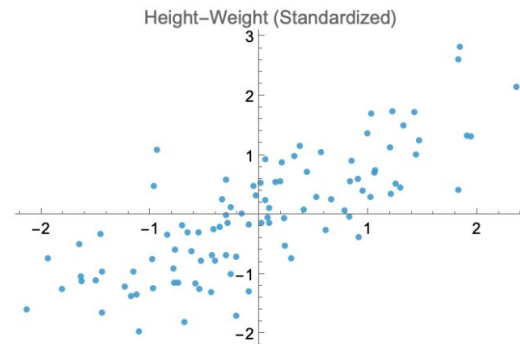
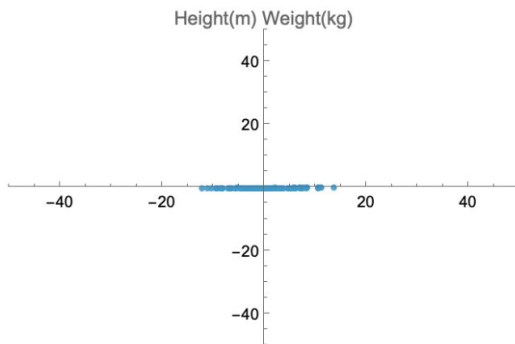
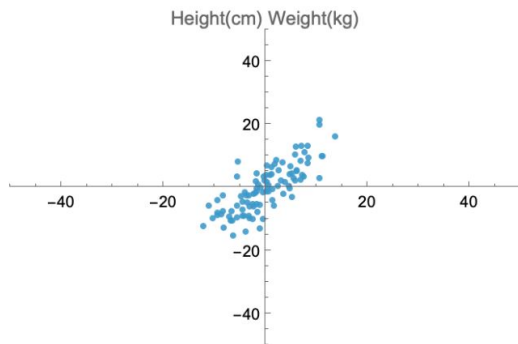
- Center the data: $X_i - \bar{X}$
- Standardize (typically): $\frac{X_i - \bar{X}}{\sigma}$
- Point cloud before/after centering





Step 1: Standardizing The Data

- Visual: why is it important to standardize?
 - Remove dependency on units
 - Get rid of scaling differences



Height vs weight of a 100 person sample



Step 2: Finding Covariance Matrix

- Find the covariance matrix:

$$\text{Cov}(\vec{X}) = \begin{bmatrix} \text{Var}(X_1) & \text{Cov}(X_1, X_2) & \cdots & \text{Cov}(X_1, X_m) \\ \text{Cov}(X_2, X_1) & \text{Var}(X_2) & \cdots & \text{Cov}(X_2, X_m) \\ \vdots & & \ddots & \vdots \\ \text{Cov}(X_m, X_1) & \text{Cov}(X_m, X_2) & \cdots & \text{Var}(X_m) \end{bmatrix}$$

- Computational shortcut: if M is the matrix of your standardized data. Then

$$\text{Cov}(\vec{X}) = \frac{1}{N} M^T M$$



Step 3: Finding Eigenvectors and Eigenvalues

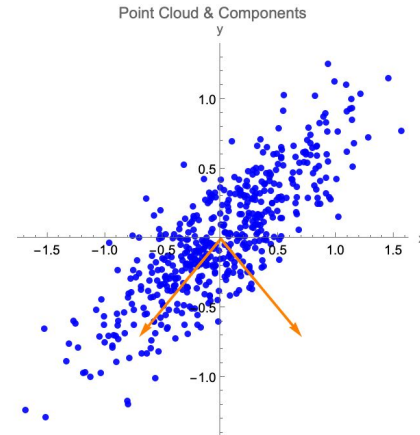
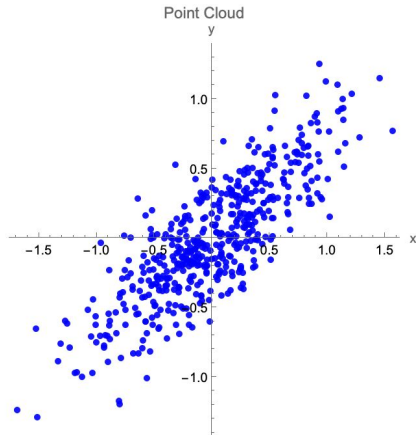
If the data is standardized, the eigenvalues of the covariance matrix $\text{Cov}(\vec{X})$ measure the proportion of the variance in the direction of the corresponding eigenvectors.

$$\sum_{i=1}^N \lambda_i = \sum_{i=1}^N \text{Var}(X_i) = N$$

The eigenvector of largest eigenvalue will determine the first principal component,
The eigenvector of second largest eigenvalue will be the second principal component,
Etc.

Step 4: Choose Number of Principal Components

- Choose a number N of principal components
- Pick the eigenvectors with the largest N eigenvalues



First two principal components



Step 4: Choosing Number of Principal Components

Deciding the number of components onto which the

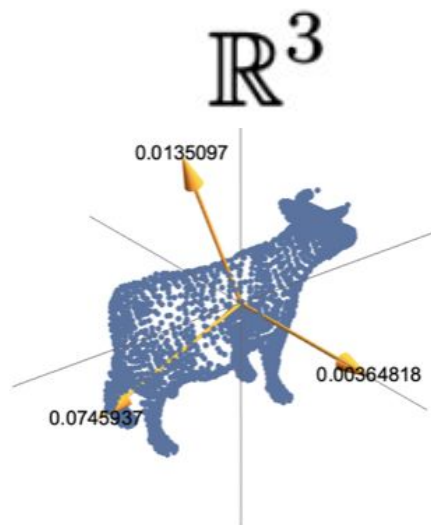
- For visualization purposes 2 or 3 (obvious reasons)
- Elbow Rule
- Scree Plot



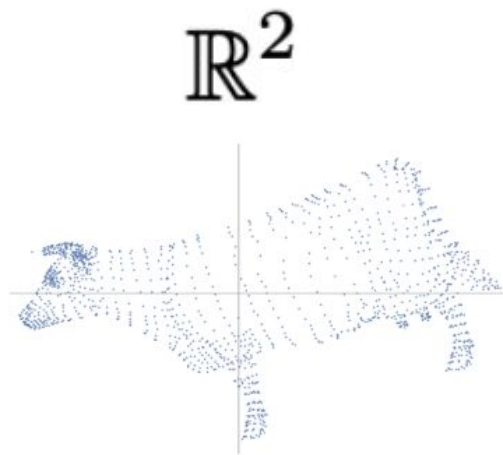
Feature Extraction

Example: From \mathbb{R}^3 to \mathbb{R}^2

Projection onto the **first two** principal components.



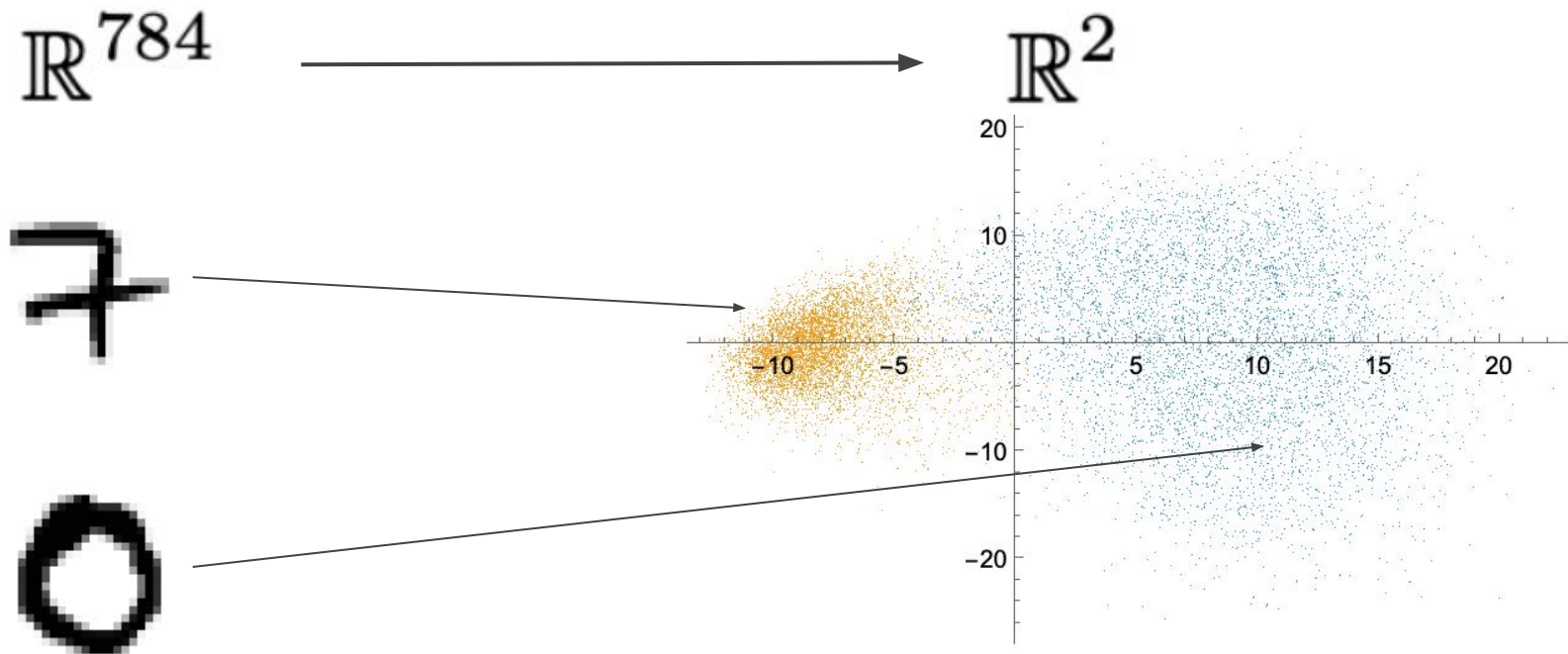
3D Cow



2D Cow



Example: MNIST Classification





Example: Medical Imaging

$$\mathbb{R}^{784} \longrightarrow \mathbb{R}^2$$



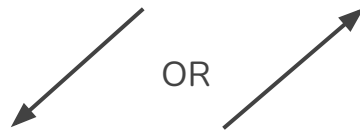
Principal Component Analysis: Summary

- Standardize (or center) each feature
- Compute covariance matrix
- Find eigenvectors and eigenvalues of the covariance matrix
 - The eigenvalues represent the proportion of overall variance in the direction of the eigenvector
 - Select a number of eigenvectors according to their eigenvalues
 - Project the data onto those eigenvectors
 - Find combinations of feature that are more relevant to the overall variance



Subtleties, Remarks, and Coding

- Built-in algorithms will center your data, but (typically) won't standardize it.
- There is a sign ambiguity when choosing the eigenvectors.
- There are some rules about how many principal components eigenvectors to choose.



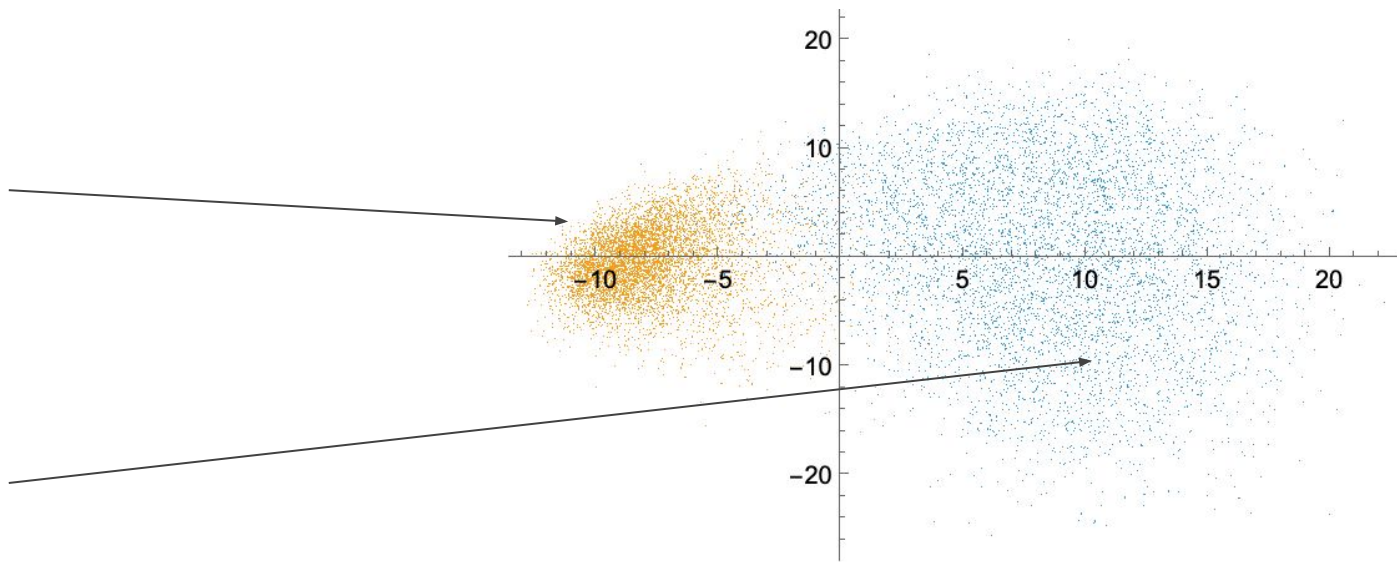
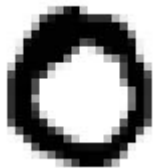
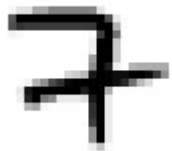


PCA + Other Algorithms





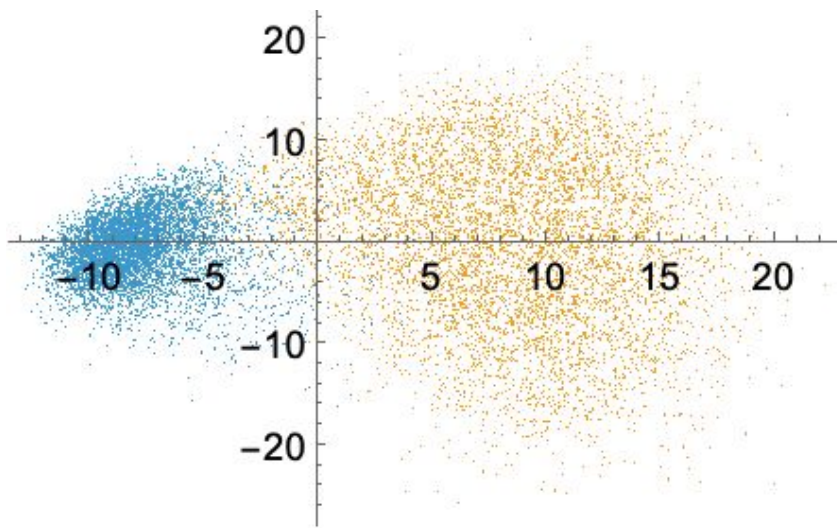
Running Example: Classifying 0s and 7s



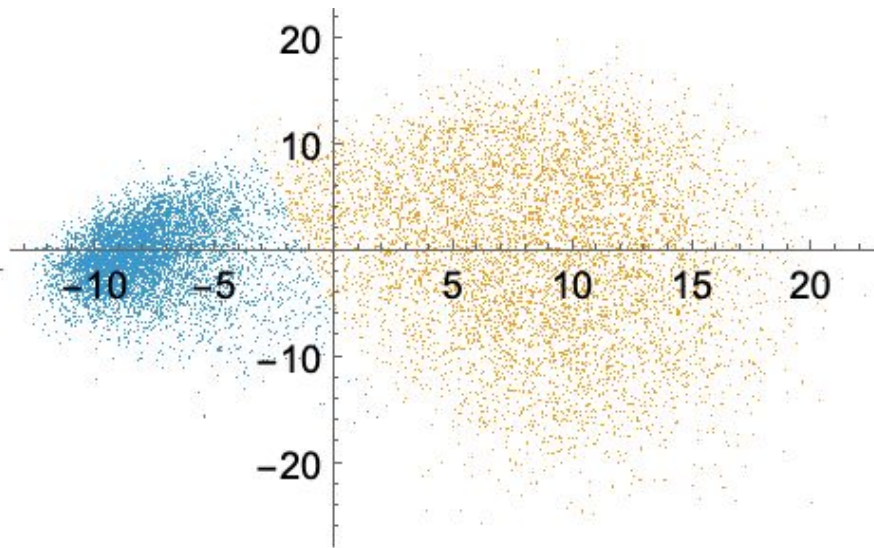


PCA + Logistic Regression

- First run Logistic Regression
- Then apply PCA
- Timing: 27s



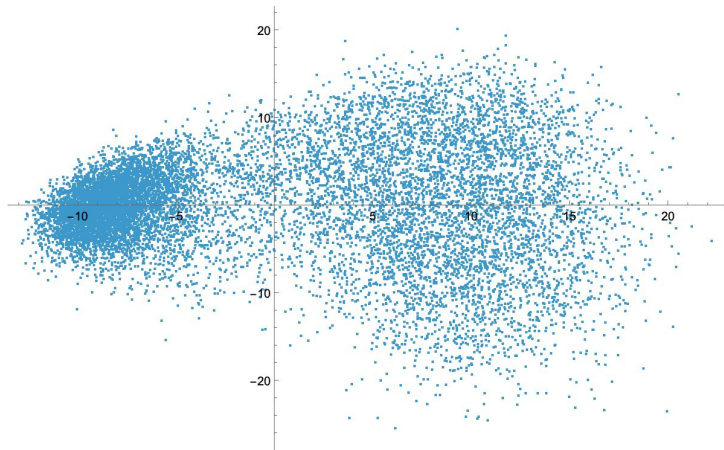
- First apply PCA
- Then run Logistic Regression
- Timing: 1.9s



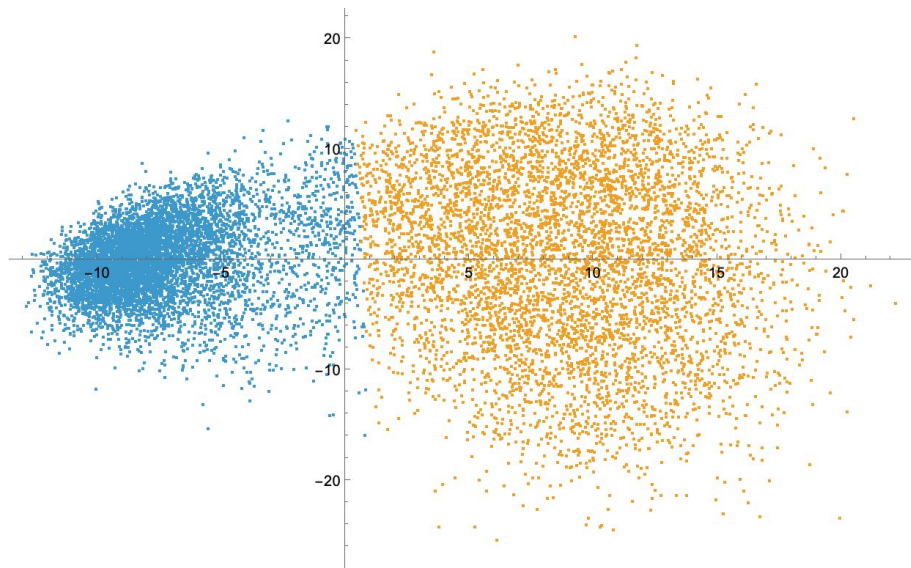


PCA + Clustering

- First find 2 clusters
- Apply PCA
- Timing: 2.6s
- Finds 1 cluster (and 1 singleton)



- First apply PCA
- Find 2 clusters
- Timing: 0.5s





Other Dimensional Reduction Algorithms

