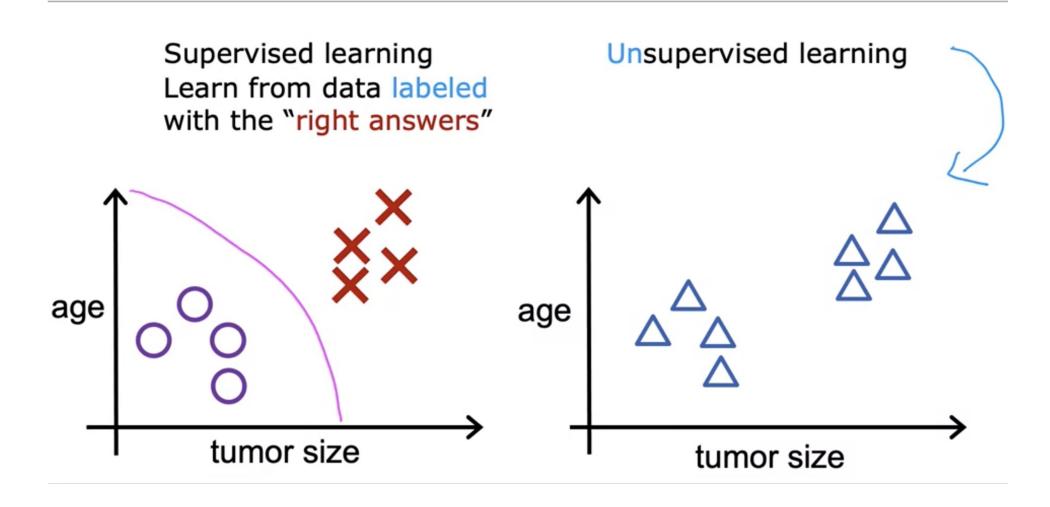




Introducing Unsupervised Learning



Unsupervised Learning

Clustering
Dimension reduction

Clustering: Google news

Giant panda gives birth to rare twin cubs at Japan's oldest zoo

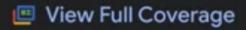
USA TODAY · 6 hours ago

- Giant panda gives birth to twin cubs at Japan's oldest zoo
 CBS News · 7 hours ago
- Giant panda gives birth to twin cubs at Tokyo's Ueno Zoo
 WHBL News · 16 hours ago
- A Joyful Surprise at Japan's Oldest Zoo: The Birth of Twin Pandas

The New York Times · 1 hour ago

Twin Panda Cubs Born at Tokyo's Ueno Zoo

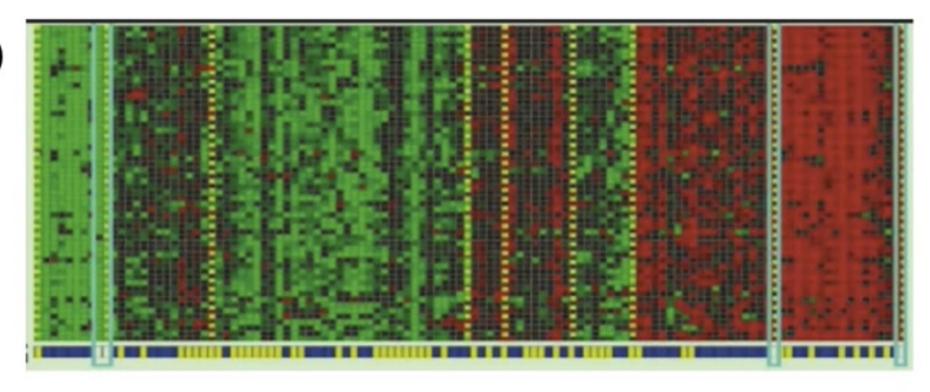
PEOPLE · 6 hours ago





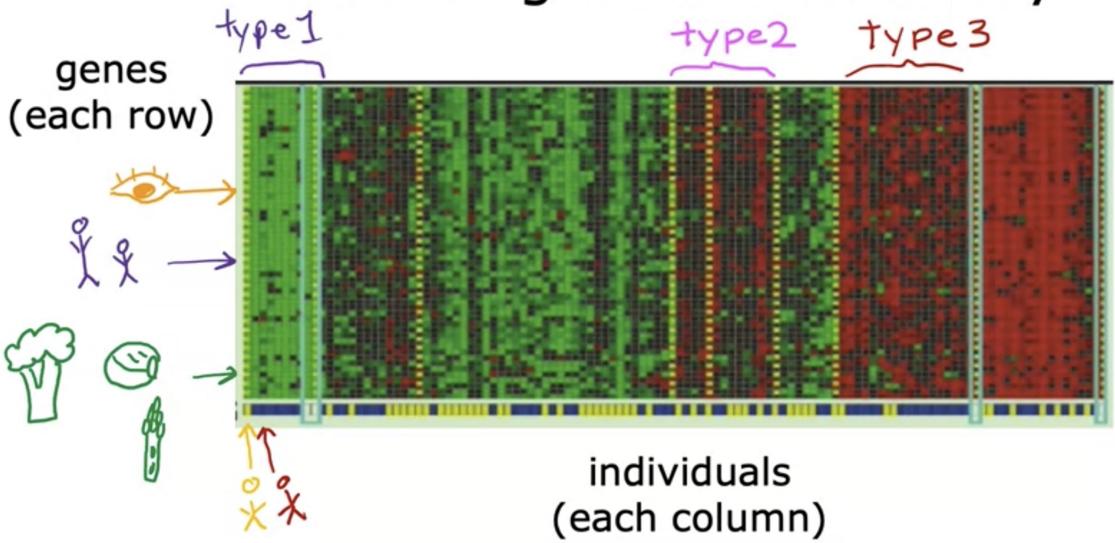
Clustering: DNA microarray

genes (each row)

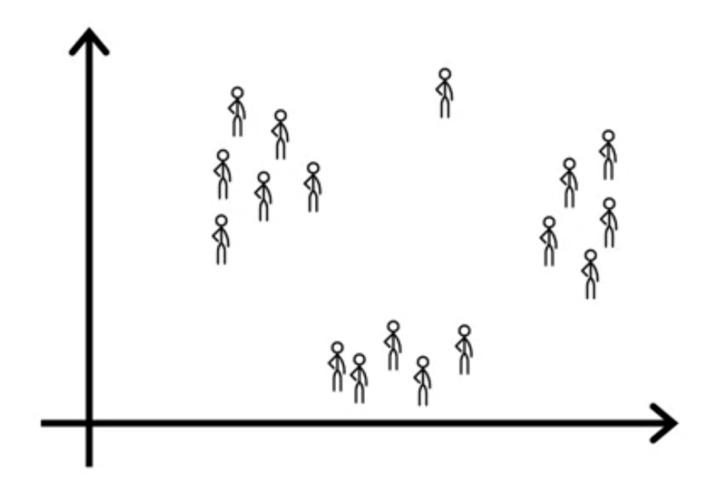


individuals (each column)

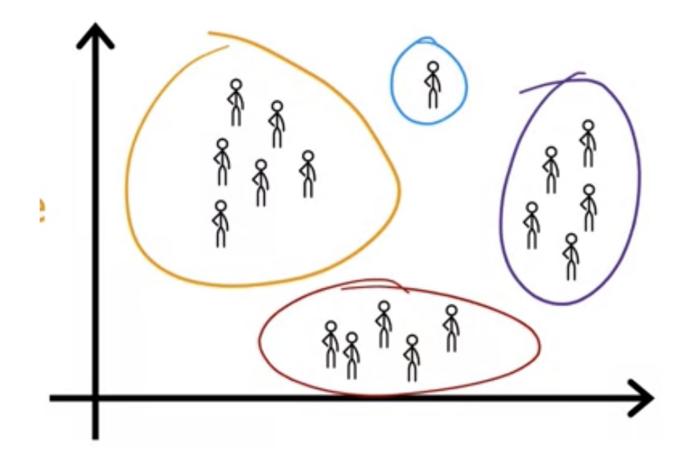
Clustering: DNA microarray



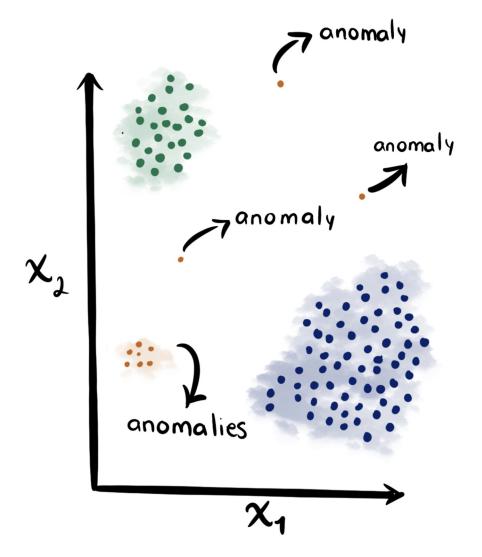
Clustering: Grouping customers



Grouping Customers



Anomaly Detection



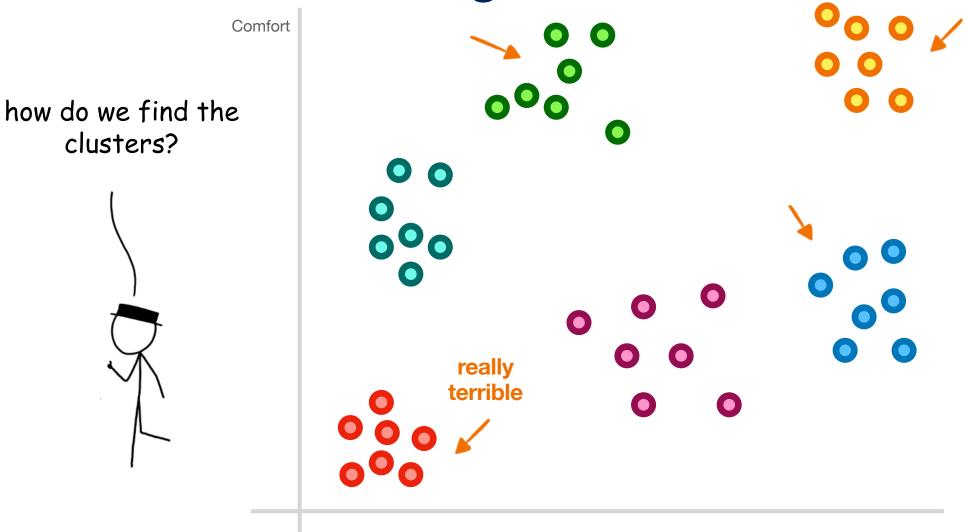
Credit: Anomaly Detection

Unsupervised Learning

Clustering

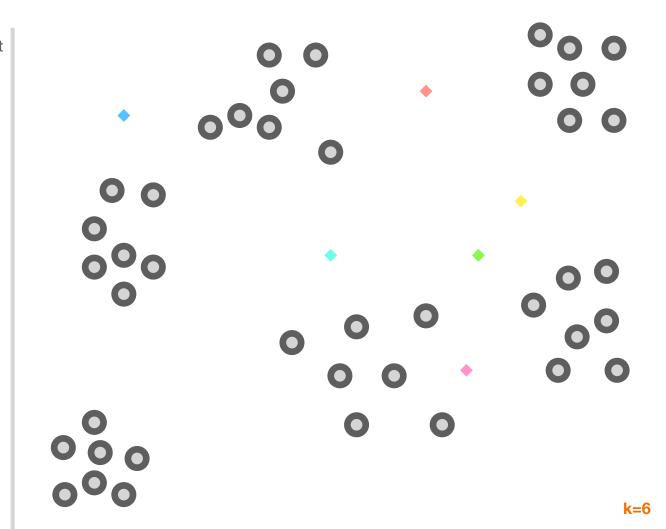
Dimension reduction

K-means clustering

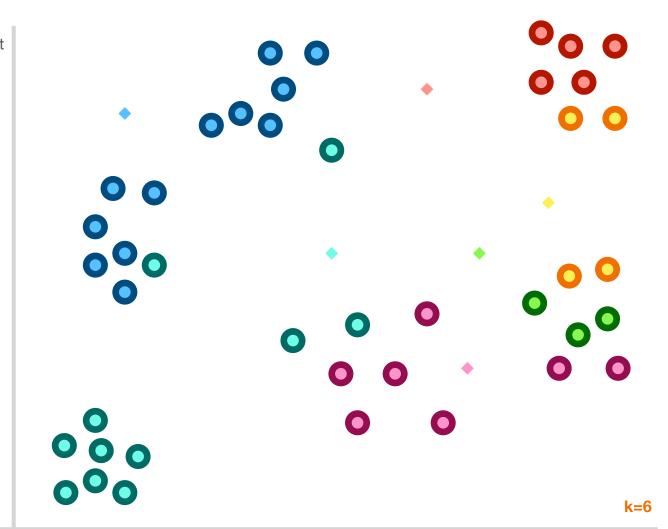


clusters can tell us specifics about the relationship of data unsupervised unsupervised learning!

- 1. pick a K-number of clusters
- 2. randomly pick a series of "centroids"
- 3. assign each particle to the **centroid** closest to it

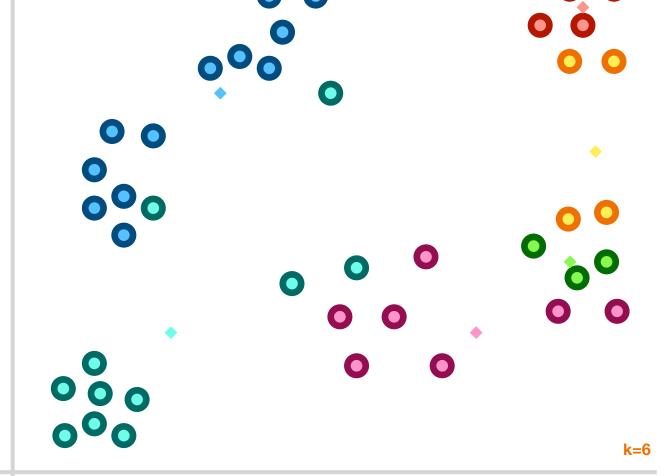


- 1. pick a K-number of clusters
- 2. randomly pick a series of "centroids"
- 3. assign each particle to the **centroid** closest to it 4.move the **centroid** to the weighted geometric center of samples assigned to it



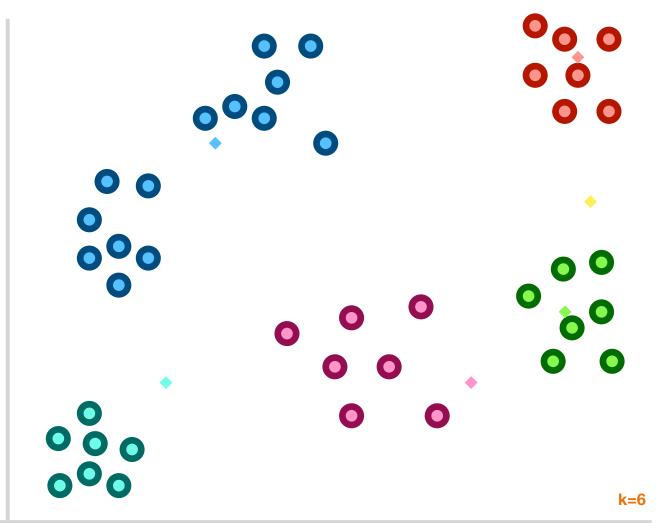
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centroids stop moving!



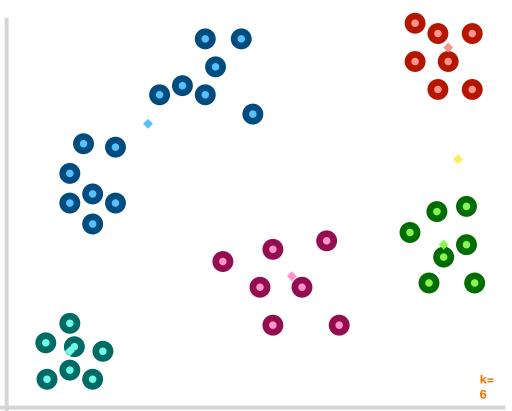
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centroids stop moving!



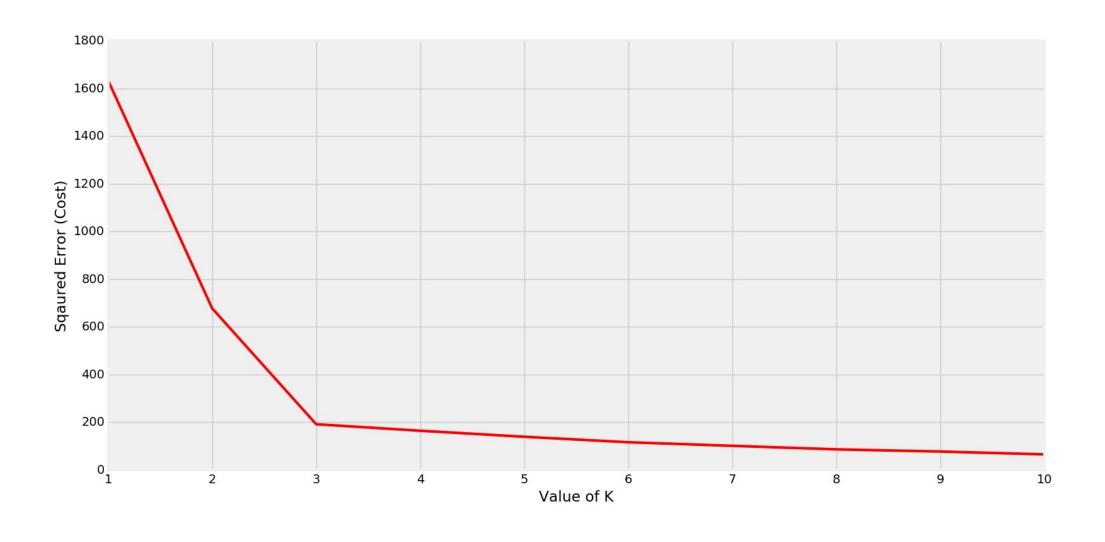
Did we get back the same clusters? Nope. And that's OK.

Fashion

Did we get back the same clusters?

Nope. And that's OK.

Evaluation and Choosing K



Unsupervised Learning

Clustering

Dimension reduction

Unsupervised Learning

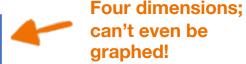
Why Dimension reduction?

Motivation for Dimension Reduction

Complex systems often must be modeled with large datasets, having dozens to millions of columns.

Often, several columns can be adding similar information to the model. So, there is a certain level of *redundancy*.

^A Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
Person A	165	65	60,000	2
Person B	168	63	100,000	5
Person C	159	82	50,000	1
Person D	183	68	90,000	4
Person E	187	87	110,000	5
Person F	189	89	95,000	4



Motivation for Dimension Reduction

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
Person A	165	65	60,000	2
Person B	168	63	100,000	5
Person C	159	82	50,000	1

What if I have a lot of features, but not a lot of samples?



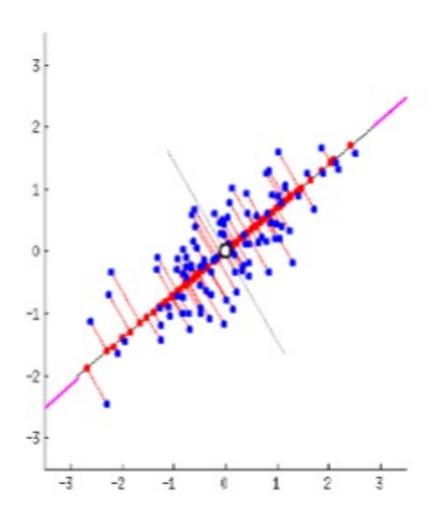
Motivation for Dimension Reduction

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
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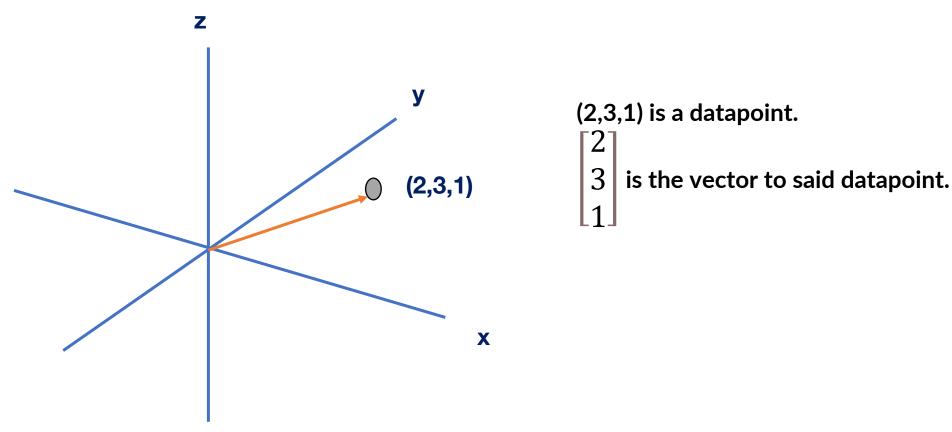
So, how do we reduce dimensionality without significant loss of information?



Enter...

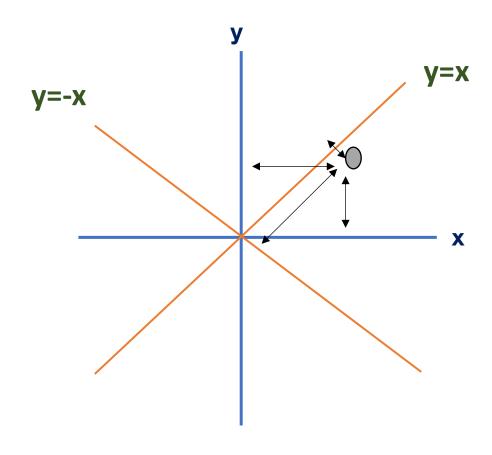


Exploring Dimensions and Basis Vectors



Dimension = # of features

Exploring Dimensions and Basis Vectors

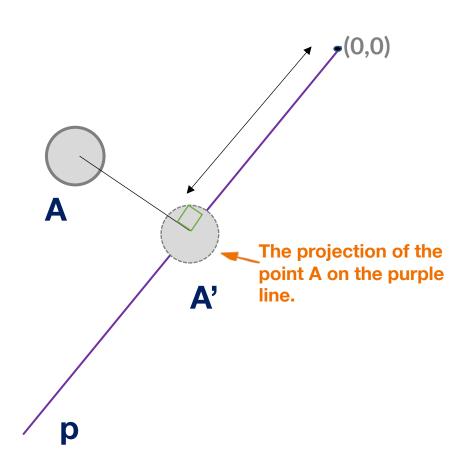


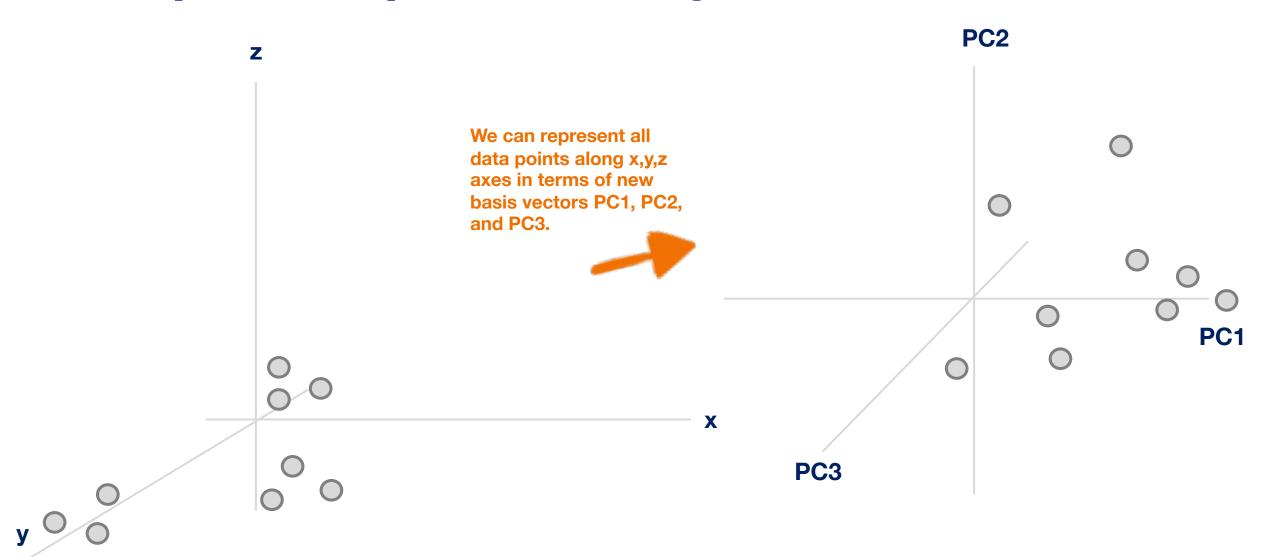
This gray point can be expressed as 3 blocks on x axis and 2 blocks on the y axis.

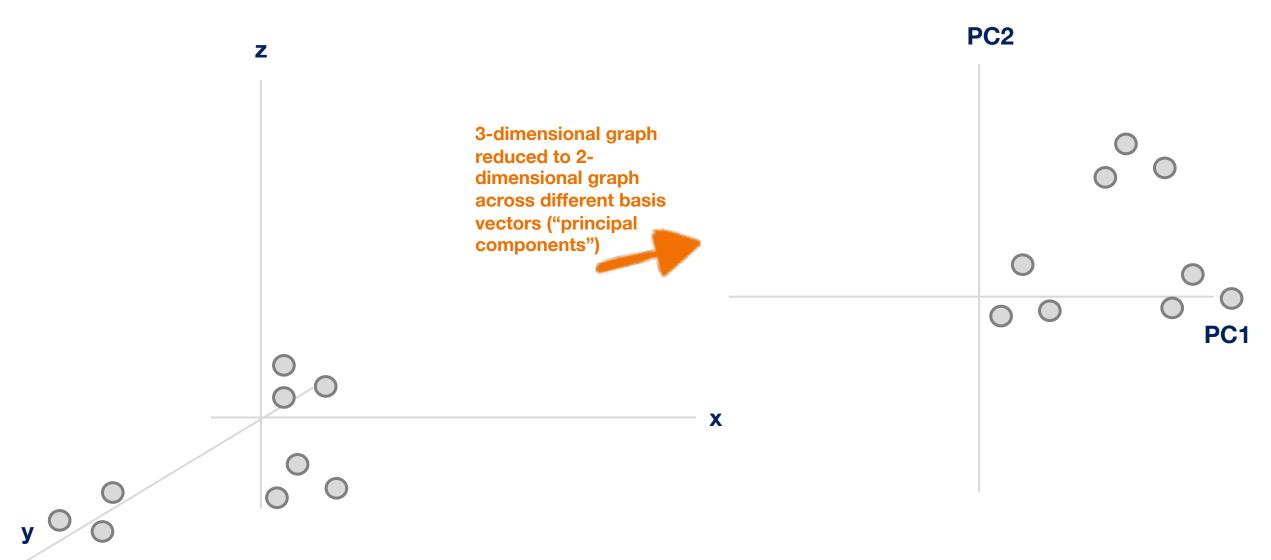
It can also be expressed as 1 block on y = -x and 3 blocks on y=x

Projection

The projection (A') of a point A on a particular line p is the point such that the line AA' is perpendicular to p.







How do we decide which PCs to drop when reducing the dimensionality of the data?



Principal Components

Think of these as new axes that we are orienting our data across.

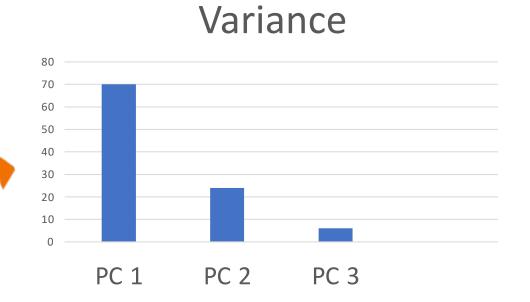
So instead of x,y, z, rather some linear combination of them.

How do we decide which PCs to remove when reducing the dimensionality of the data?

They are done such that each principal component is uncorrelated with the others, so that translation across each component indicates different information. **So, they represent directions of maximal variance.**

This allows differences between data points to become more prominent

Represents percentage of variance for each PC. Notice how PC1 has the most and it drops after that.

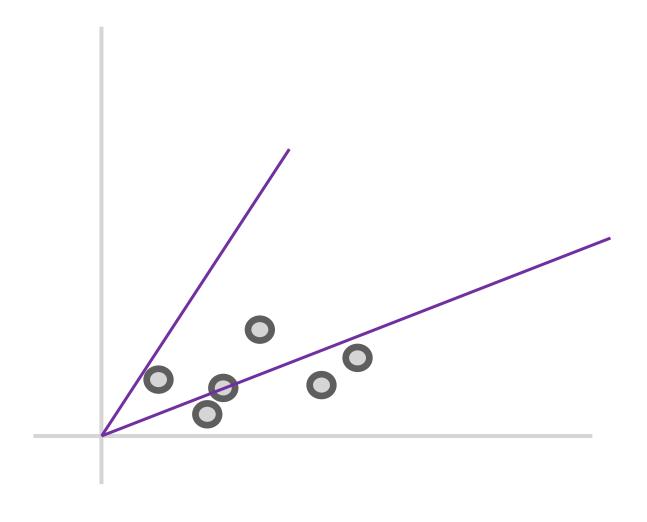


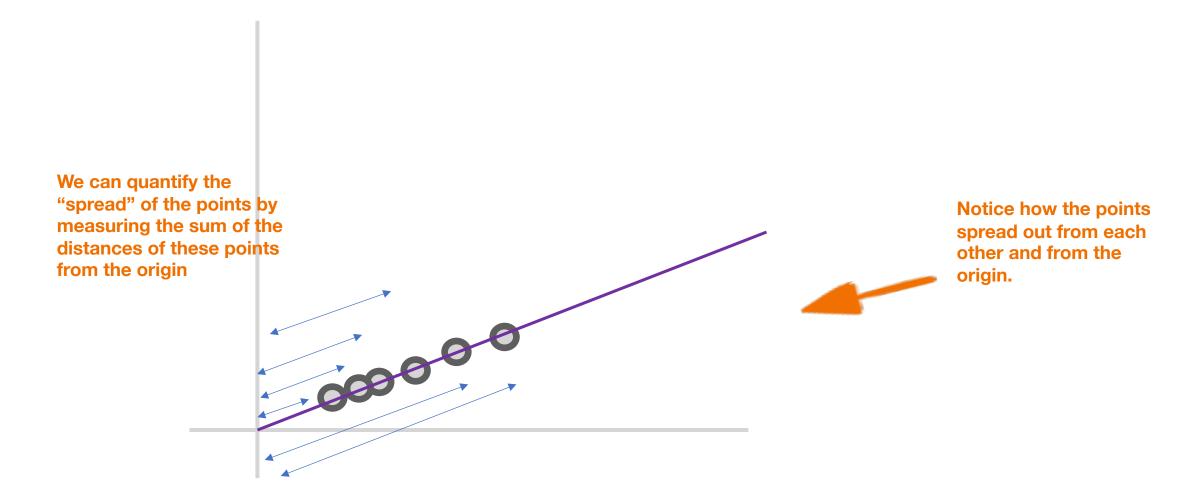


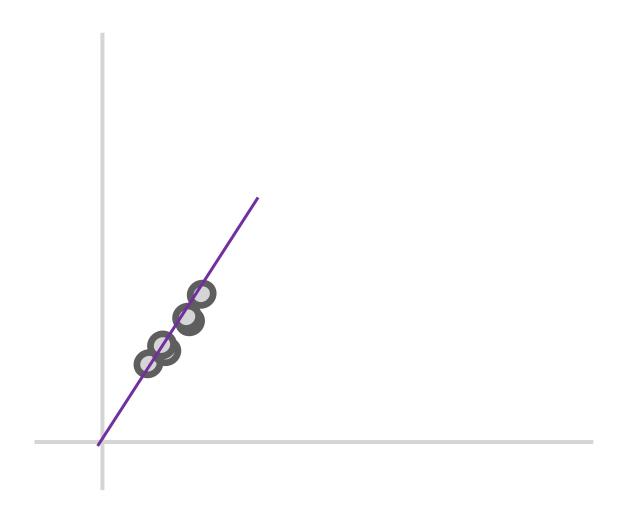
Since PC3 accounts for a very small percentage of overall variance, we can remove it. This is how PCA reduces dimensionality

How do we decide the PCs?





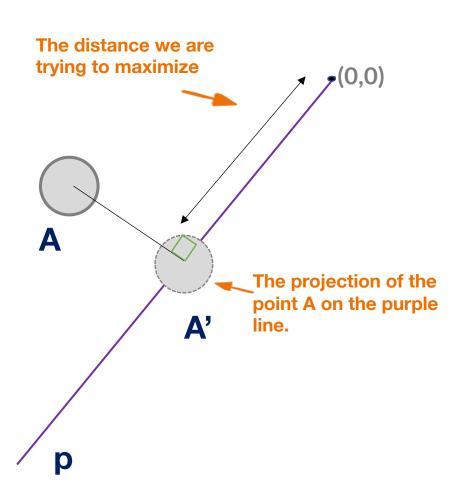




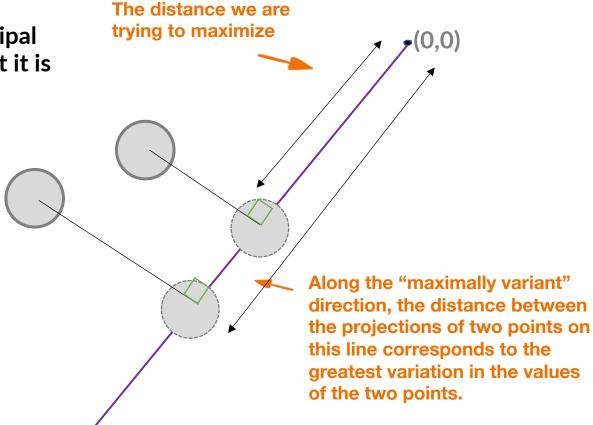
We can quantify the "spread" of the points by measuring the sum of the distances of these points from the origin

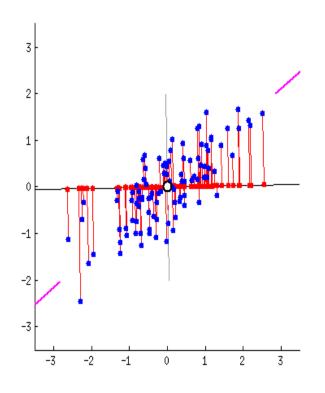
1st base or "Principal Component 1". Line that maximizes sum of distances of projections of points from origin. In essence, maximizes variance of distribution.

The projection (A') of a point A on a particular line p is the point such that the line AA' is perpendicular to p.



Idea behind this principal component line is that it is an axis along the "maximally variant" direction.

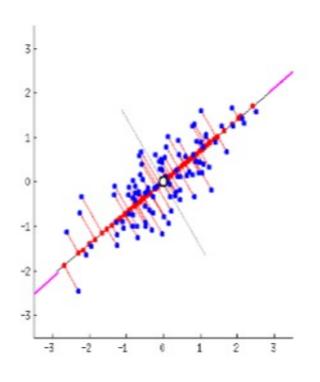




Maximizing the variance along the line

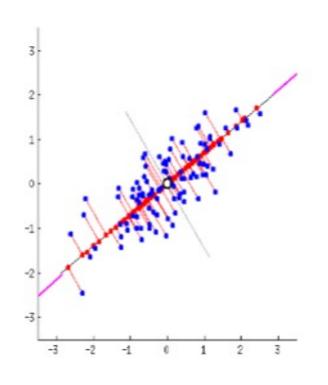


Built using https://gist.github.com/anonymous/7d888663c6ec679ea654287 15b99bfdd



Maximizing the variance along the line







Standardization

Covariance Matrix Calculation

Eigenvector Calculation

Form Principal Components and Build Graph

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
Person A	165	65	60,000	2
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Compare the data of each of the 4 columns. How do they differ numerically?

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
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Person F	189	89	95,000	4
Range	159-189	63-89	50,000-110,000	1-5
Variance	161.76	135.87	564166666	2.7

Compare the data of each of the 4 columns. How do they differ numerically?

Their range varies drastically. Consequently, their variances are very different.

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
Person A	165	65	60,000	2
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Person C	159	82	50,000	1
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Person F	189	89	95,000	4
Range	159-189	63-89	50k-100k	1-5
Variance	161.76	135.87	564166670	2.7

If this is not addressed, some of the feature columns will **dominate** over the other ones.

This can bias the results and final principal component analysis; making it difficult to view differences between values in one column compared to another.

So final graph may have the differences between the weights of various persons be miniscule.

So, how do we adjust our data so these differences are not as drastic?



Recap: we want to put different variables on the same scale.

This can mean many things from giving them the same mean and standard deviation, to keeping the range consistent, and so on.

Here, we will use a method called **z-scoring**.

$$z = \frac{value - mean}{standard\ deviation}$$

The rescaled distribution will have a mean of 0 and standard deviation of 1

Note: it does not mean the new data follow Normal distribution

Standardization

Covariance Matrix Calculation

Eigenvector Calculation

Form Principal Components and Build Graph

Covariance Matrix Calculation

Covariance is really just a measure of how correlated two variables/features are.

If your covariance is positive, that means there's a positive correlation.

If your covariance is positive, that means there's a negative correlation.

$$Cov(x,y) = \sum \frac{(x_i - \bar{x})(y_i - \bar{y})}{N-1}$$

Covariance Matrix Calculation

Review: Lecture 7; feature engineering

Make new features with high variance.

Pick new features with low correlation to other features.

What should our new features look like?



Covariance Matrix Calculation

Can measure this correlation using **covariance**. If covariance is **positive**, then features are correlated in the sense they both increase together. If covariance is **negative**, then features are inversely correlated.

$$\begin{bmatrix} Cov(x,x) & Cov(x,y) & Cov(x,z) \\ Cov(y,x) & Cov(y,y) & Cov(y,z) \\ Cov(z,x) & Cov(z,y) & Cov(z,z) \end{bmatrix}$$

$$Cov(x,y) = \sum \frac{(x_i - \bar{x})(y_i - \bar{y})}{N-1}$$

Standardization

Covariance Matrix Calculation

Eigenvector Calculation

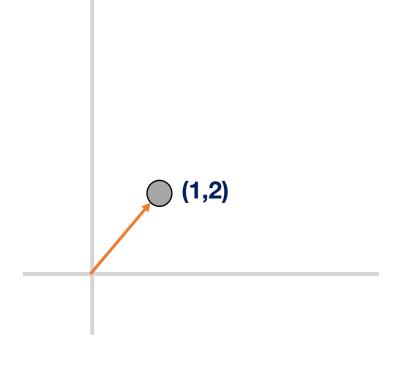
Form Principal Components and Build Graph

We can think of matrices as **transformations** of vectors.

When you multiply a matrix with a vector; two things happen:

- 1. It **scales** the vector.
- 2. It **rotates** the vector

$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = \begin{bmatrix} 5 \\ 4 \end{bmatrix}$$

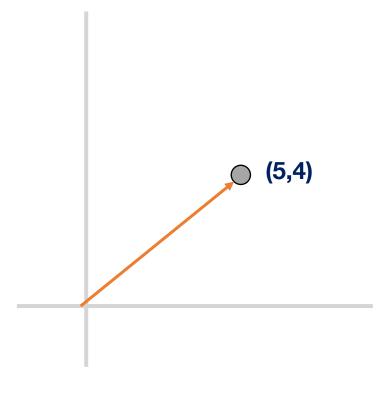


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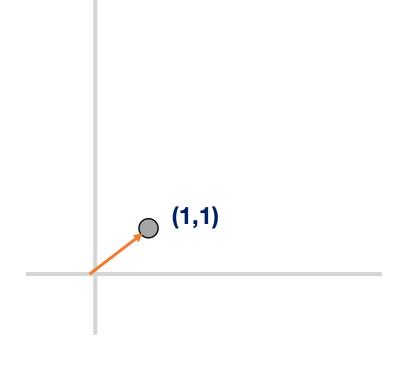
$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = \begin{bmatrix} 5 \\ 4 \end{bmatrix}$$



Eigenvectors are characteristic vectors specific to a matrix or transformation.

Graphically speaking, when you multiply a matrix with its specific eigenvectors, the eigenvectors don't get rotated, only scaled.

$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 3 \\ 3 \end{bmatrix} = 3 \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

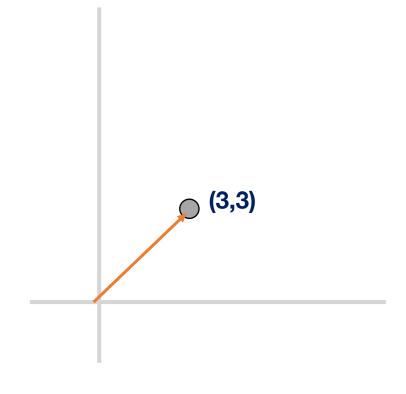


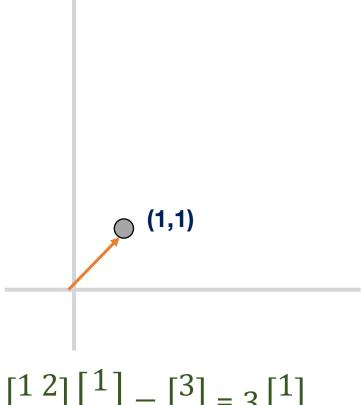
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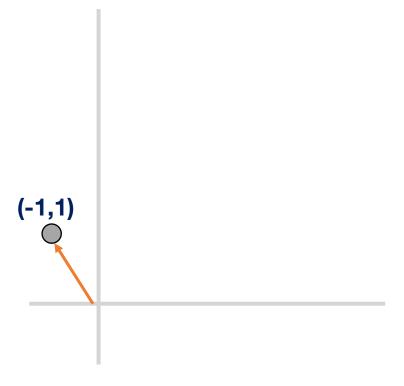
The factor by which an eigenvector is scaled is called its eigenvalue

$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 3 \\ 3 \end{bmatrix} = 3 \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

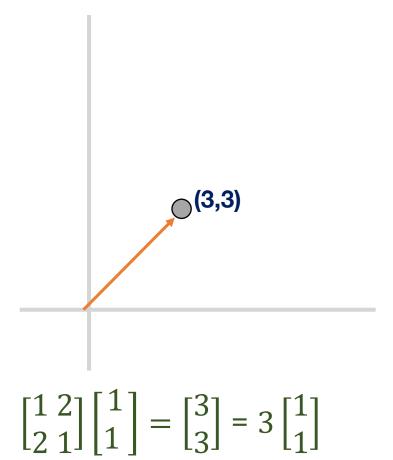


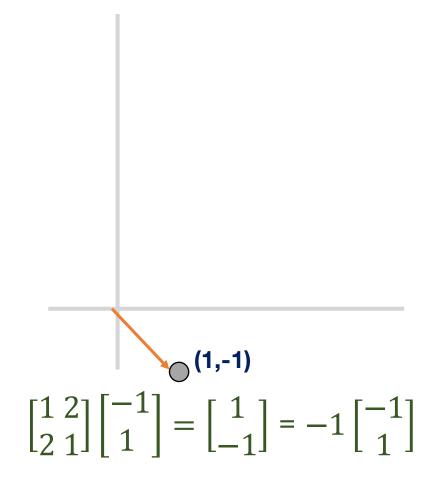


$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 3 \\ 3 \end{bmatrix} = 3 \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

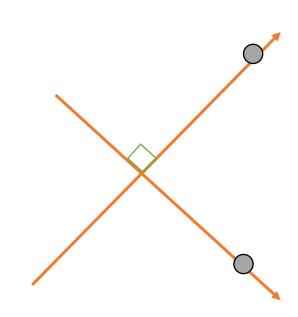


$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} -1 \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ -1 \end{bmatrix} = -1 \begin{bmatrix} -1 \\ 1 \end{bmatrix}$$



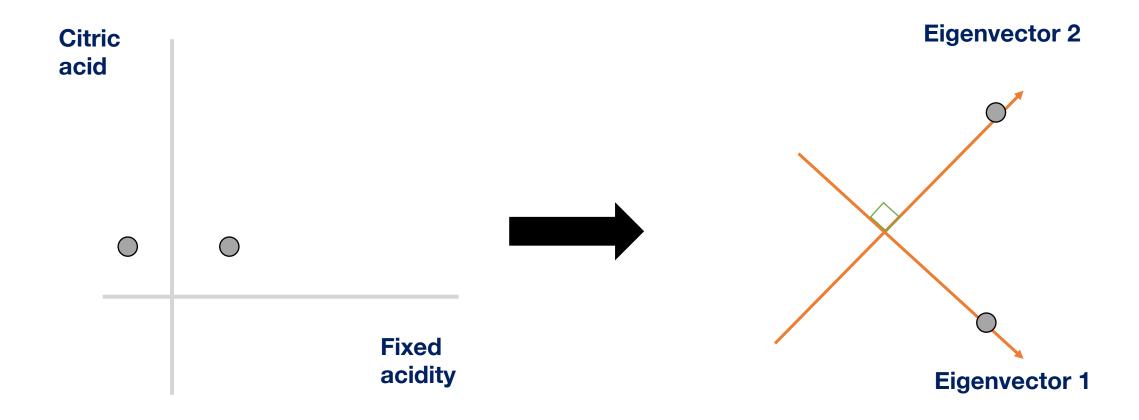


The two eigenvectors are perpendicular to each other!



Eigenvectors act as basis vectors!

Every point in 2-D can be expressed as some combination of (1,1) and (-1,1).



What matrix do we find the eigenvectors of to get our "new features" in PCA?



By calculating the eigenvectors of the covariance matrix, we can get our **principal** components.

We use the eigenvectors to create a basis for the graph. These basis vectors represent the principal components.

Since these are eigenvectors of the **covariance matrix**, they represent **directions of maximal variance**.

$$A = \begin{bmatrix} Cov(x,x) & Cov(x,y) & Cov(x,z) \\ Cov(y,x) & Cov(y,y) & Cov(y,z) \\ Cov(z,x) & Cov(z,y) & Cov(z,z) \end{bmatrix}$$

$$Av = \lambda v$$

v is the eigenvector and lambda is the eigenvalue

$$Av = \lambda v$$

$$Av - \lambda v = 0$$
$$(A - \lambda)v = 0$$
$$|A - \lambda| = 0$$

$$A = \begin{bmatrix} Cov(x,x) & Cov(x,y) & Cov(x,z) \\ Cov(y,x) & Cov(y,y) & Cov(y,z) \\ Cov(z,x) & Cov(z,y) & Cov(z,z) \end{bmatrix}$$

When you find the root of the resulting polynomial, you will find all the possible eigenvalues. For each eigenvalue, plug it into the original equation to find the corresponding eigenvector v.

Standardization

Covariance Matrix Calculation

Eigenvector Calculation

Form Principal Components and Build Graph

Form Principal Components and Build Graph

Let the three eigenvalues of the three eigenvectors v_1 , v_2 , v_3 be λ_1 , λ_2 , λ_3 such that $\lambda_1 >= \lambda_2 >= \lambda_3$

Then, the principal components will be v_1 , v_2 , v_3 and the variances they carry are in the ratio of λ_1 , λ_2 , λ_3

But if the eigenvectors are from the covariance matrix which represents the correlation of all the features, where will we be removing features?



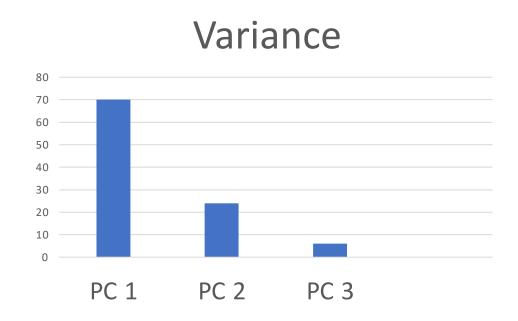
Form Principal Components and Build Graph

But if the eigenvectors are from the covariance matrix which represents the where will we be removing features?



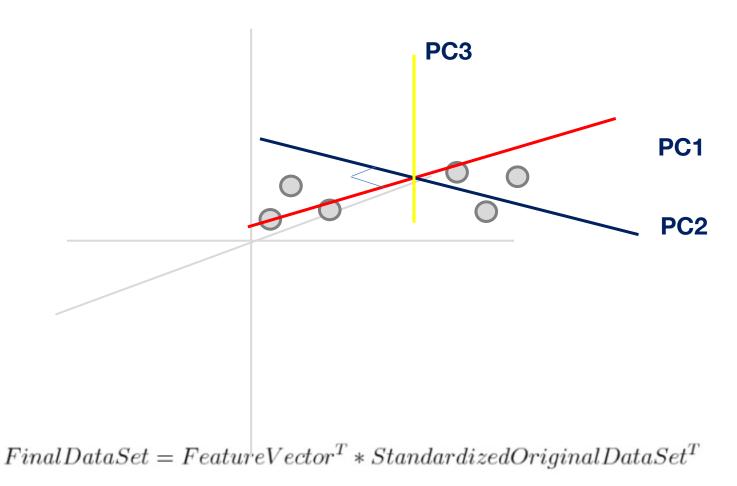
If the percentage of variance of a particular principal component is small enough, discard it. You've now removed a dimension! Form a new matrix which only has the correlation of all the features, eigenvectors/principal components you've selected.

Let this matrix be called your **Feature Vector**.

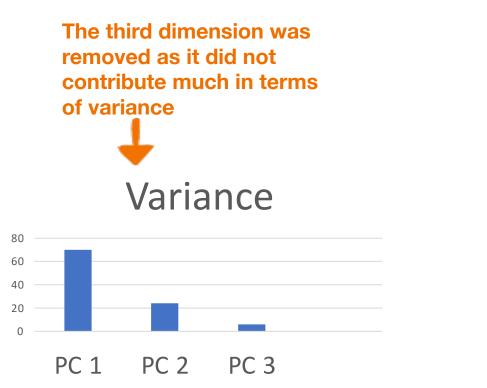


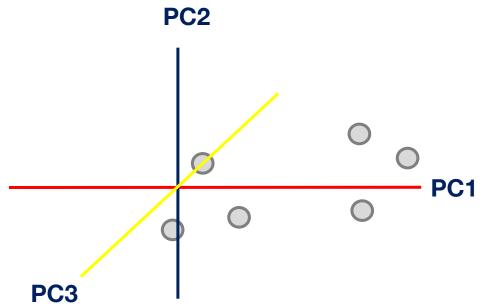
Now, it's time to reorient the original data along these new axes

Form Principal Components and Build Graph



Form Principal Components and Build Graph





 $Final Data Set = Feature Vector^T * Standardized Original Data Set^T$

Linear transformation of existing features

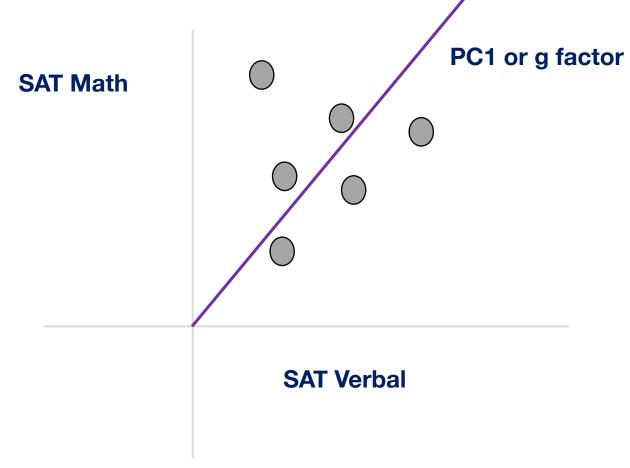
Each PC is selected to maximize the explained variance in this direction of the remain data

Dimensionality reduction is to remove dimensions with low variances

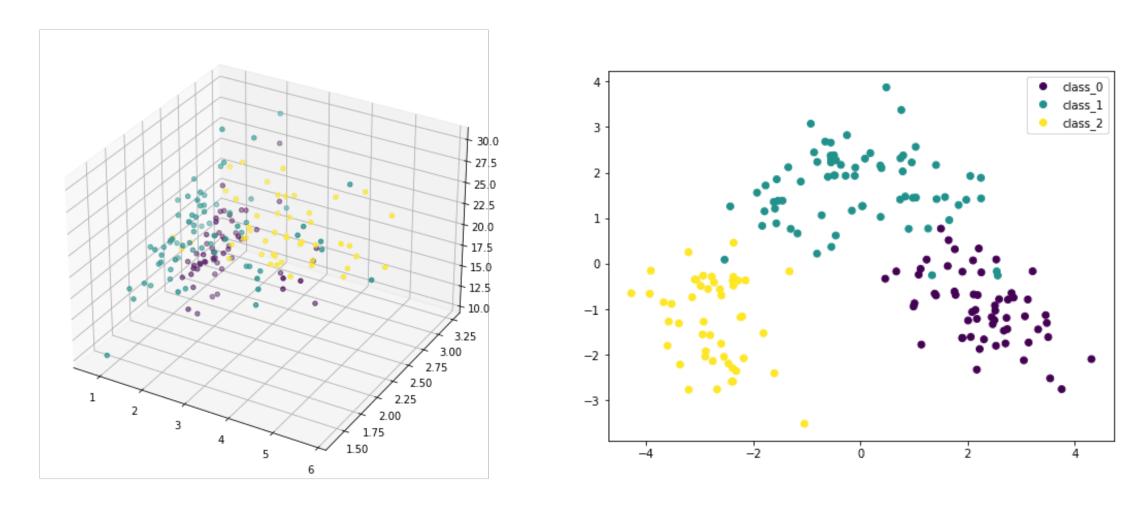
Tradeoff between dimensionality vs. info. loss

A real-world application of PCA

IQ testing!

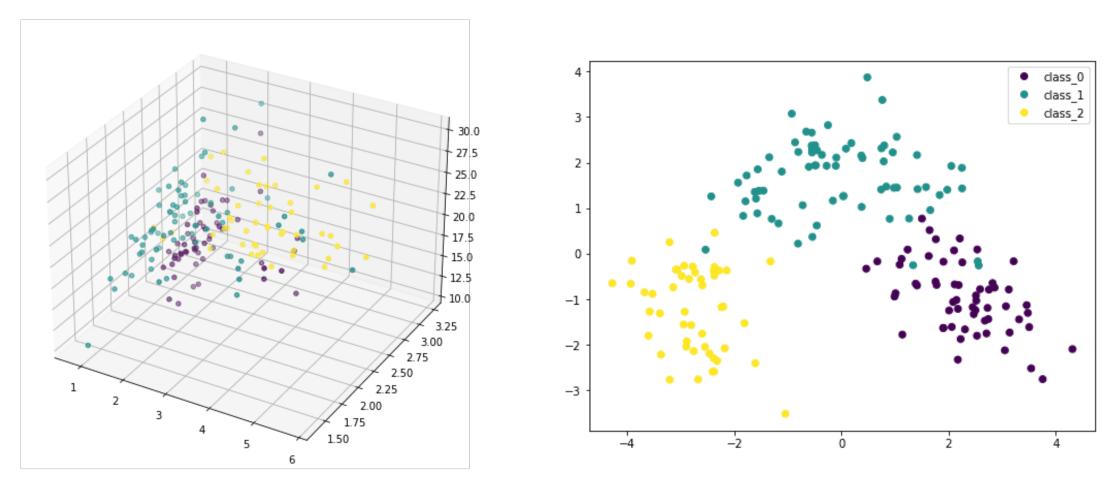


Data Visualization



The <u>wine dataset</u> with 13 features and 3 classes. Source: https://machinelearningmastery.com/principal-component-analysis-for-visualization/

Feature Extraction



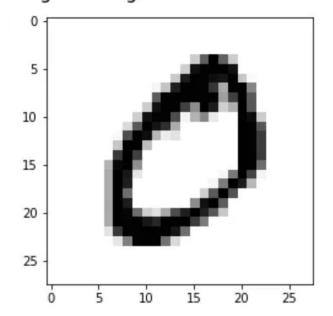
Q: feature extraction vs. feature selection?

The wine dataset with 13 features and 3 classes.

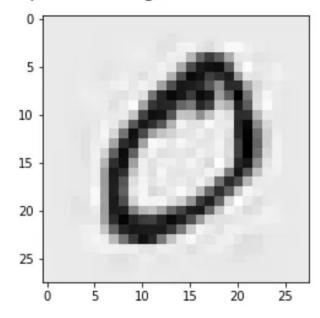
Source: https://machinelearningmastery.com/principal-component-analysis-for-visualization/

Image Compression

Original image with 784 dimensions

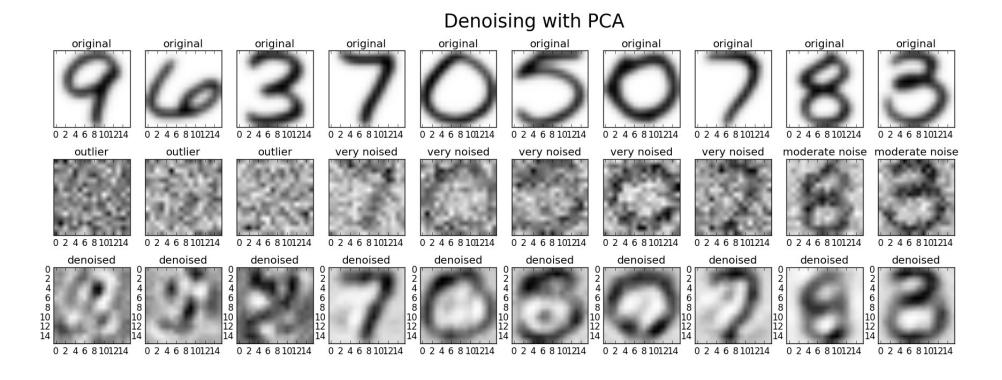


Compressed image with 184 dimensions



Source: https://towardsdatascience.com/image-compression-using-principal-component-analysis-pca-253f26740a9f

Noise Reduction



Source: https://stats.stackexchange.com/questions/247260/principal-component-analysis-eliminate-noise-in-the-data

Unsupervised Learning

Clustering
Dimension reduction

