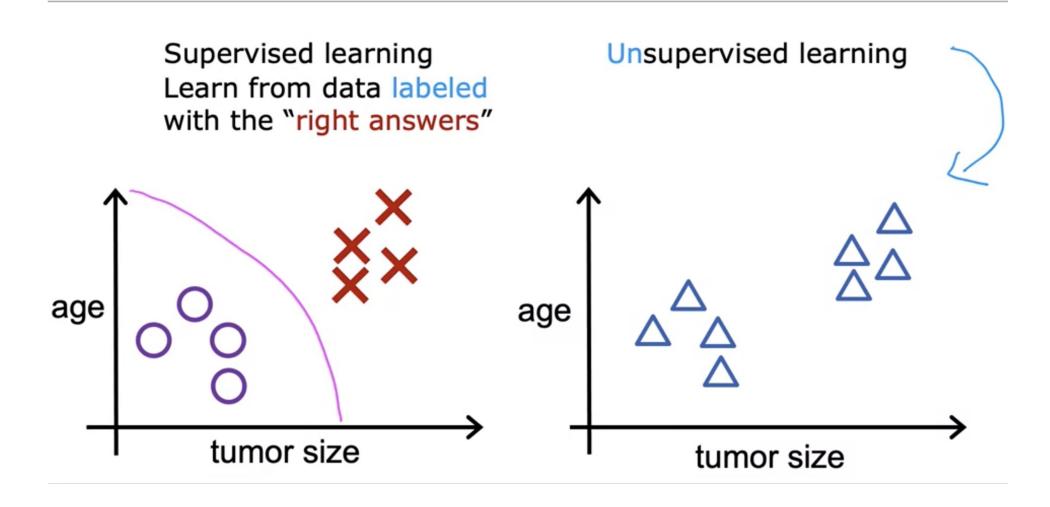




#### **Introducing Unsupervised Learning**



### **Unsupervised Learning**

Clustering
Dimension reduction

# Clustering: Google news

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

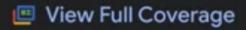
USA TODAY · 6 hours ago

- 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

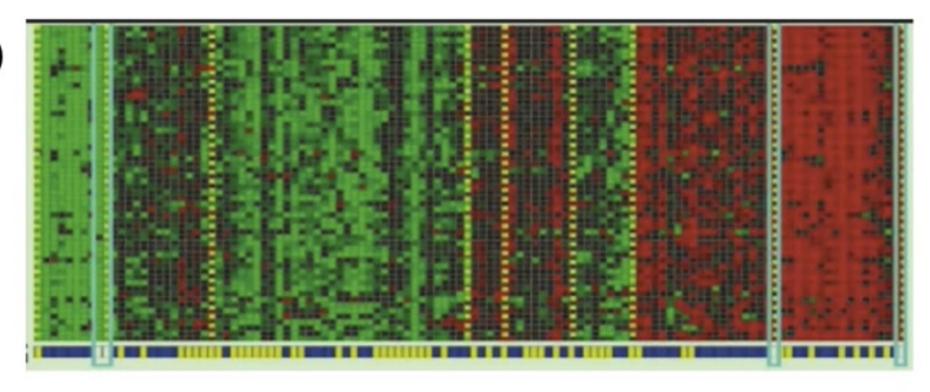
PEOPLE · 6 hours ago





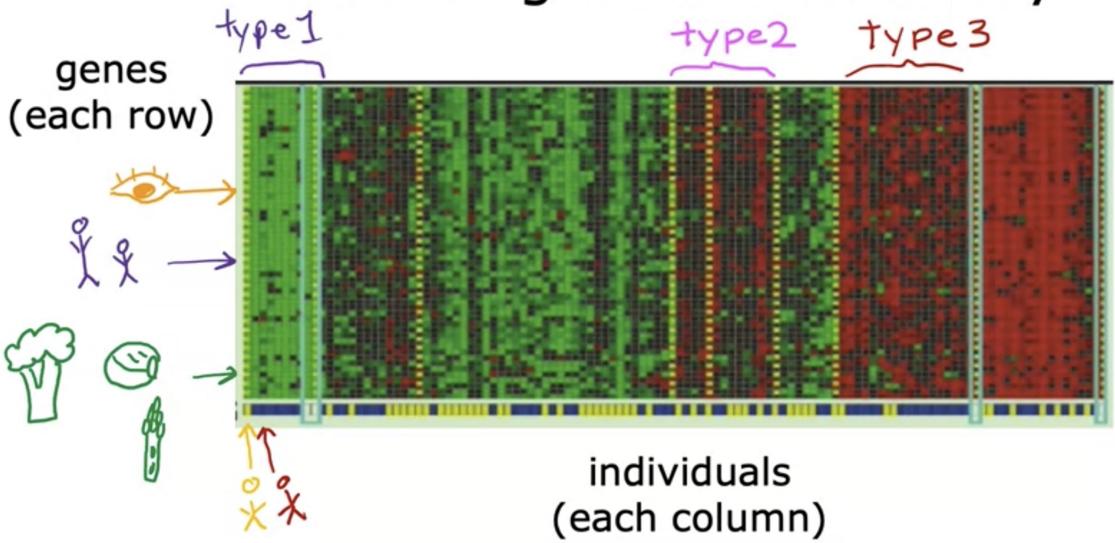
## Clustering: DNA microarray

genes (each row)

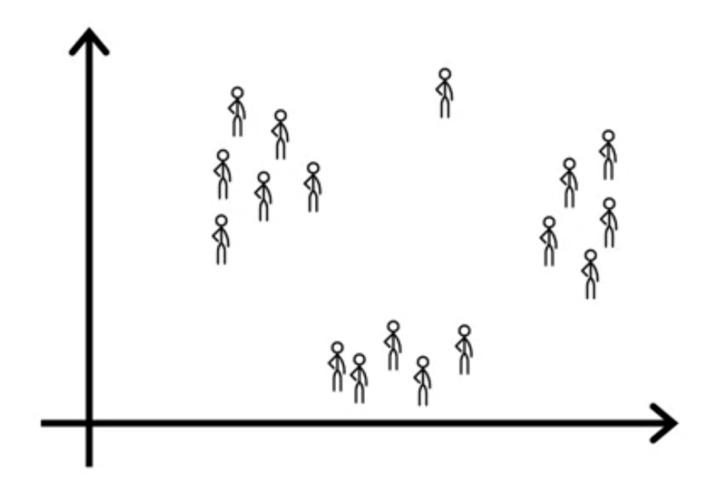


individuals (each column)

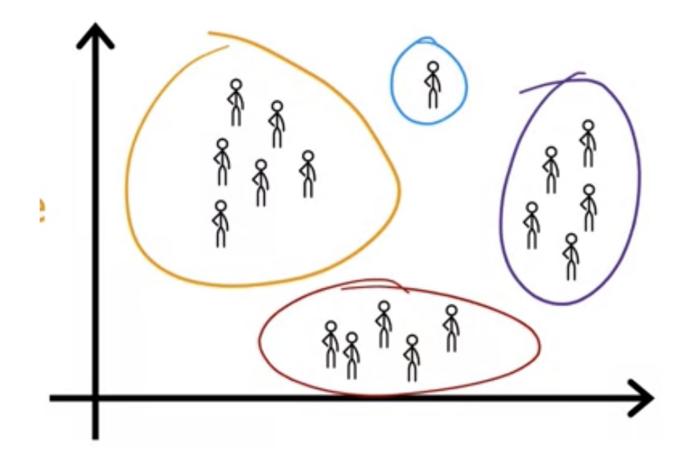
# Clustering: DNA microarray



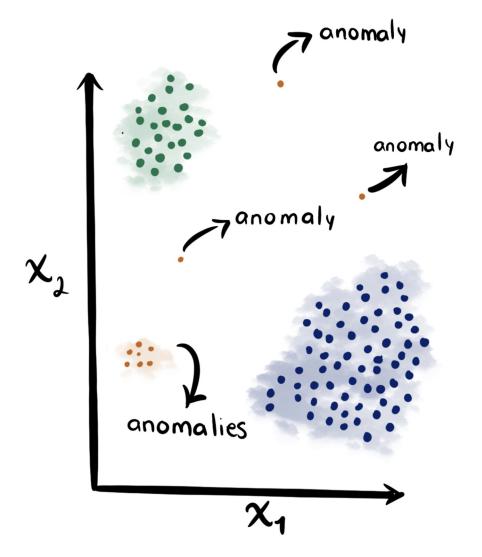
# Clustering: Grouping customers



### **Grouping Customers**

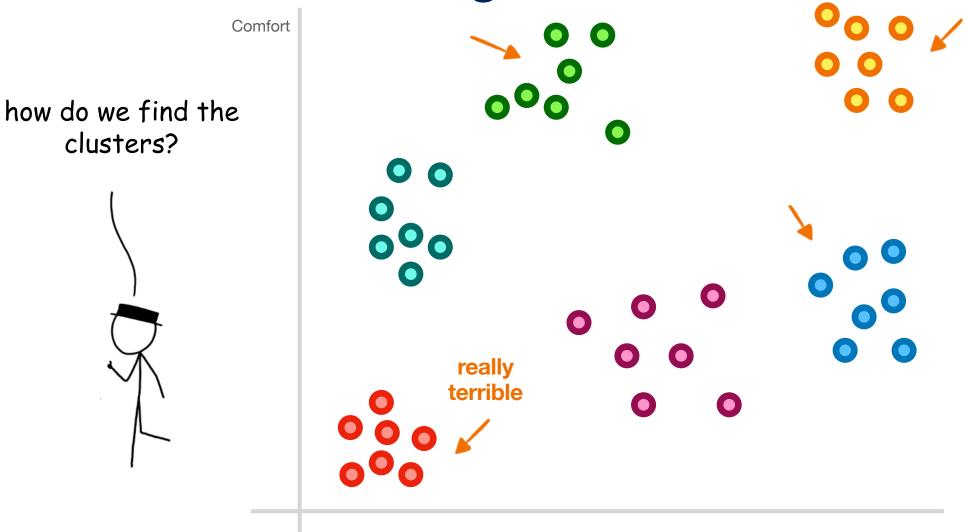


### **Anomaly Detection**



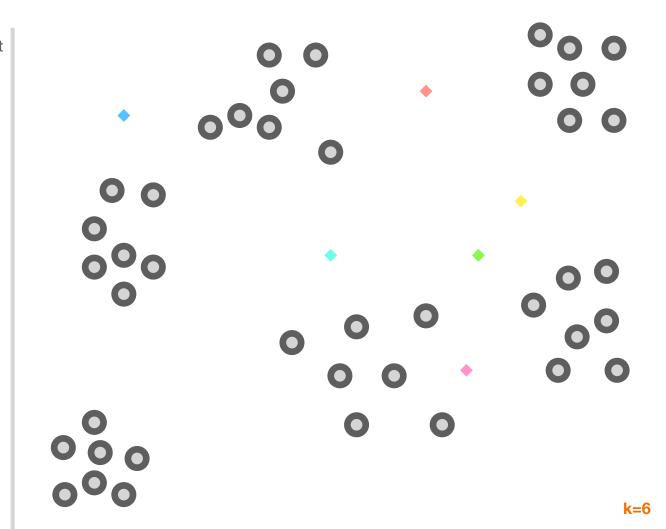
**Credit:** Anomaly Detection

### K-means clustering

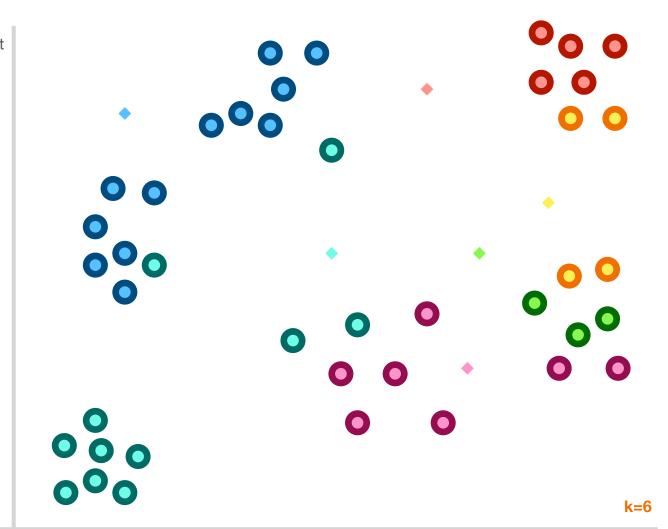


clusters can tell us specifics about the relationship of data unsupervised unsupervised learning!

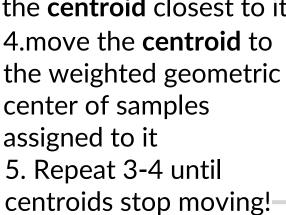
- 1. pick a K-number of clusters
- 2. randomly pick a series of "centroids"
- 3. assign each particle to the **centroid** closest to it



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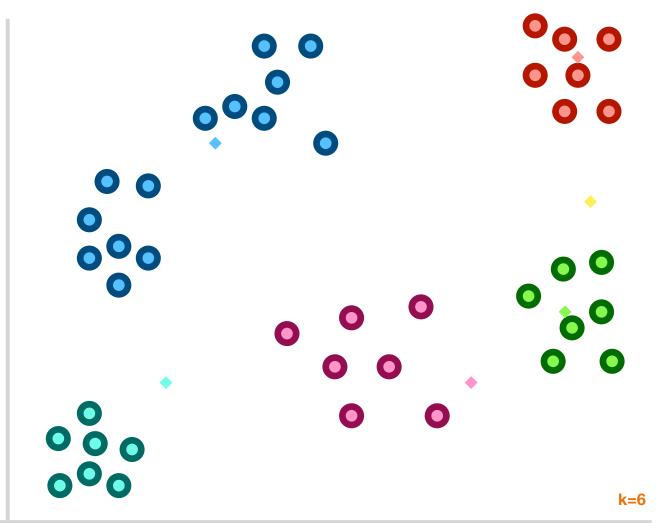
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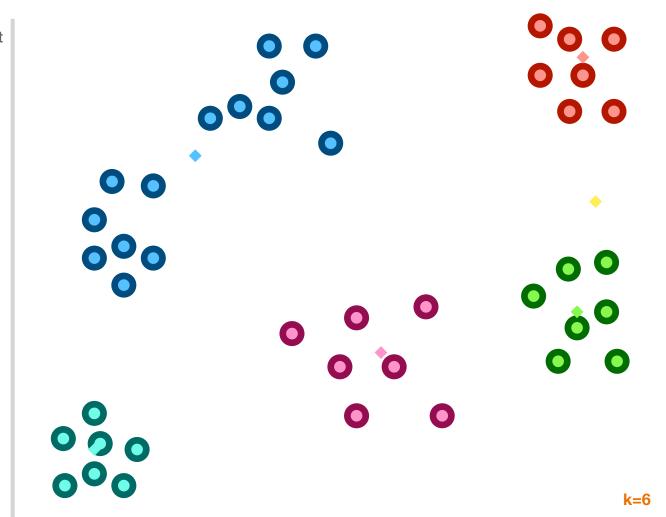
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centroids stop moving!



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centroids stop moving!



Did we get back the same clusters?

Fashion

Nope. And that's OK.

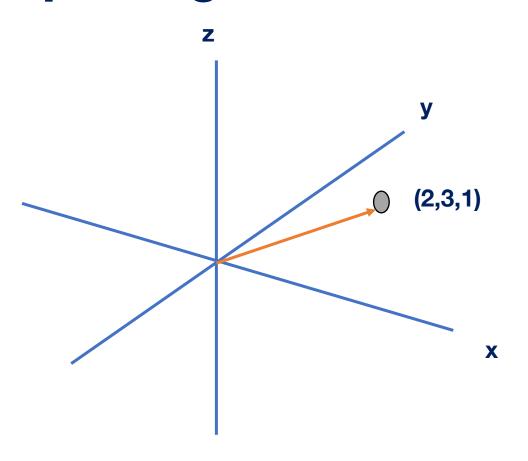
Did we get back the same clusters?

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### **Unsupervised Learning**

Clustering
Dimension reduction

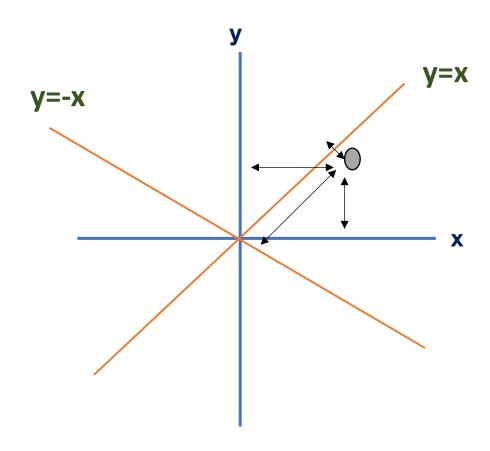
#### **Exploring Dimensions and Basis Vectors**



(2,3,1) is a datapoint.

3 is the vector to said datapoint.

#### **Exploring Dimensions and Basis Vectors**



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

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

#### **Motivation for Dimension Reduction**

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

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

Additionally, datasets with too many features may be difficult to represent graphically.

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



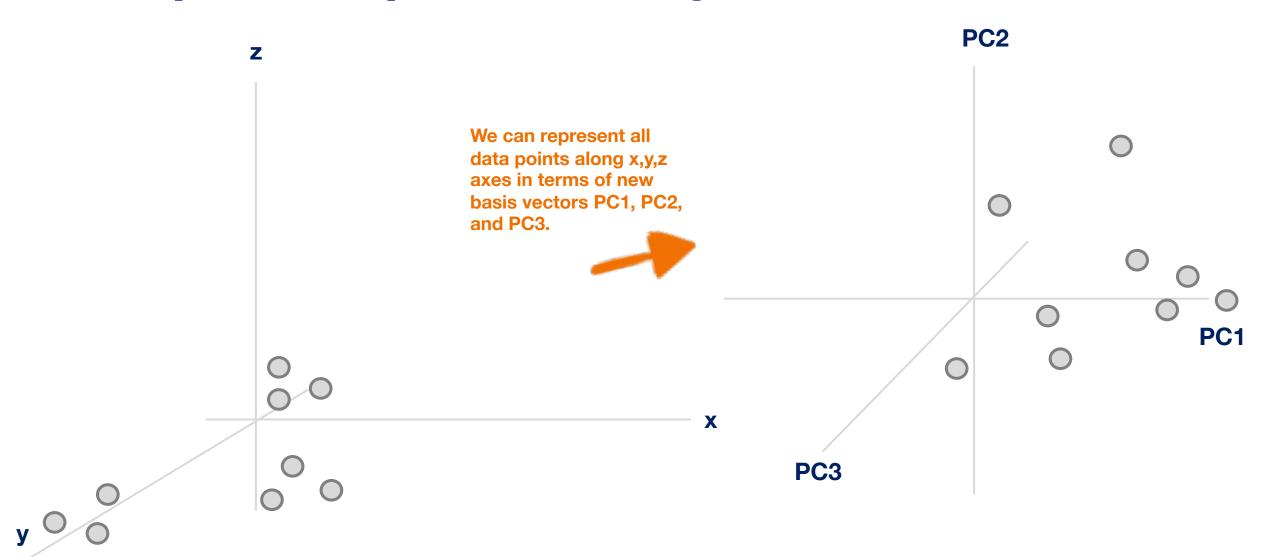
#### **Motivation for Dimension Reduction**

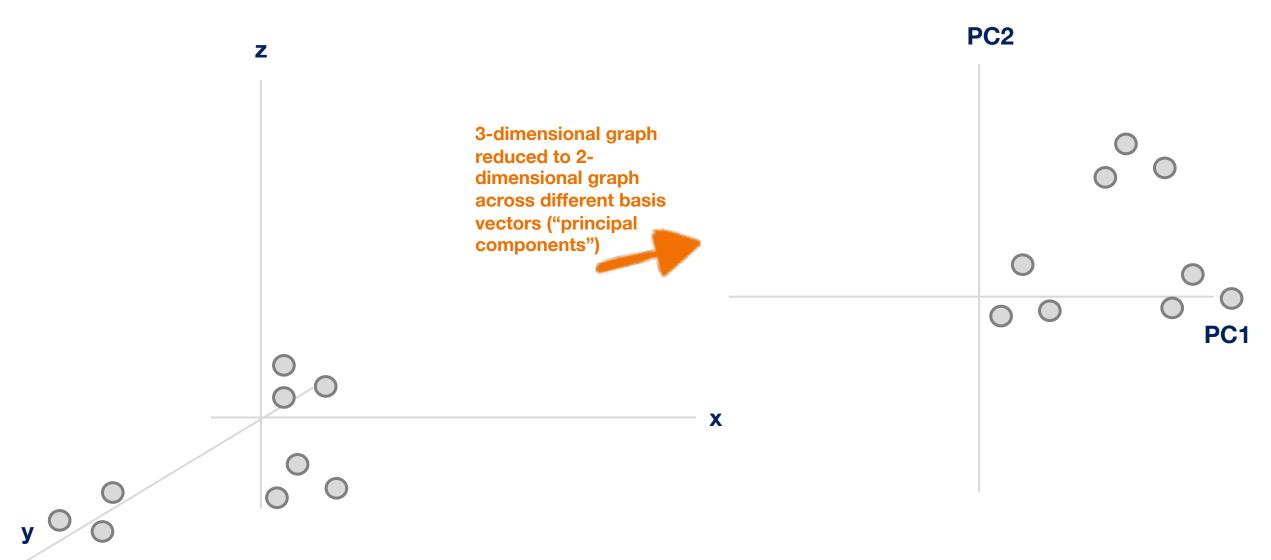
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So, how do we reduce dimensionality without significant loss of information?



Enter...





How do we decide what features to remove when reducing the dimensionality of the data?

#### **Principal Components**

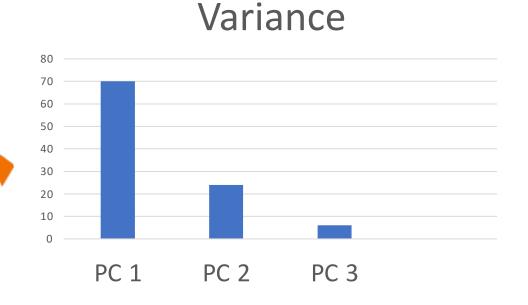
Think of these as new axes that we are orienting our data across.

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

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

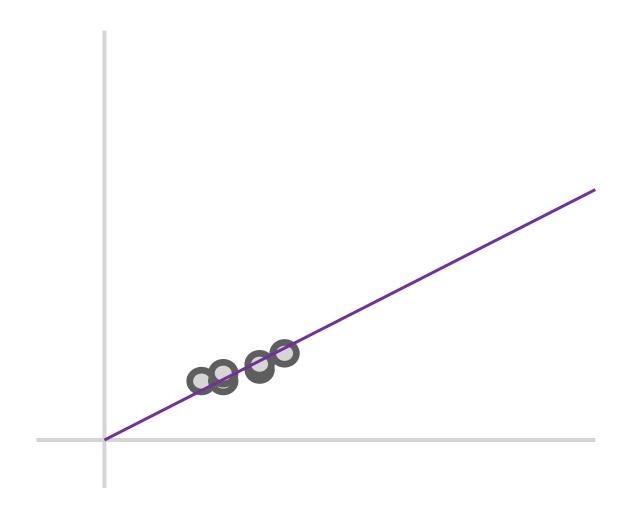
Represents percentage of variance for each PC. Notice how PC1 has the most and it drops after that.

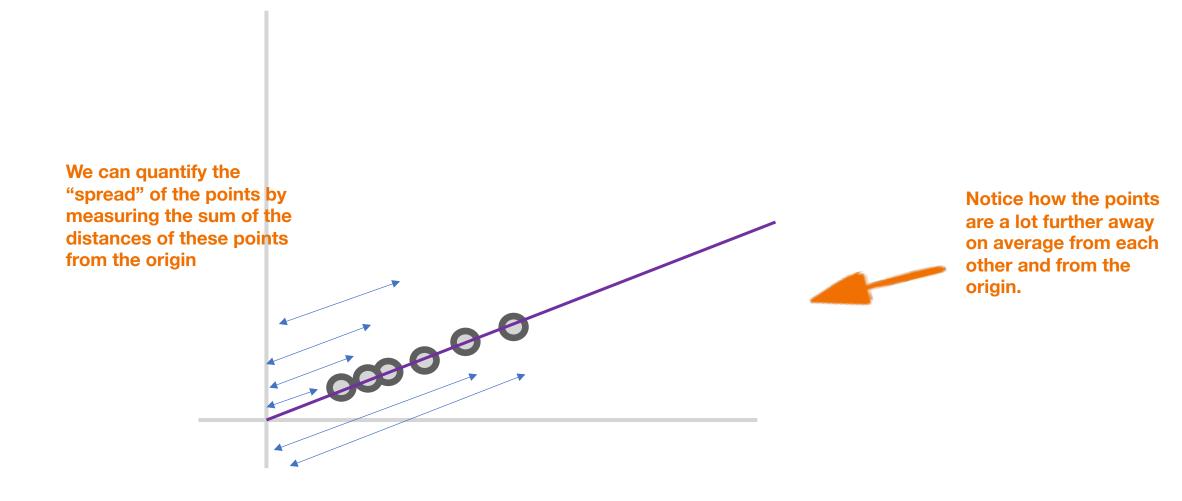


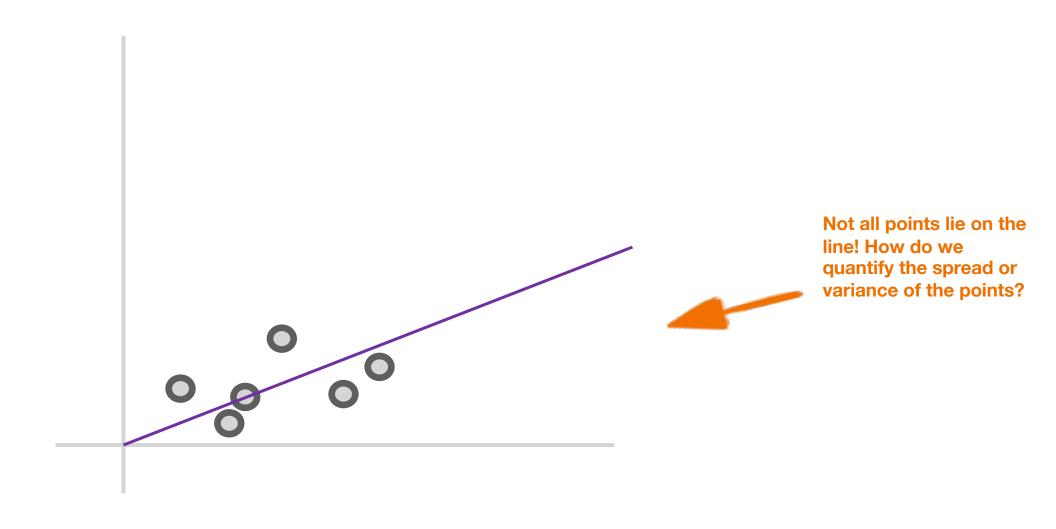
How do we decide what features to remove when reducing the dimensionality of

the data?

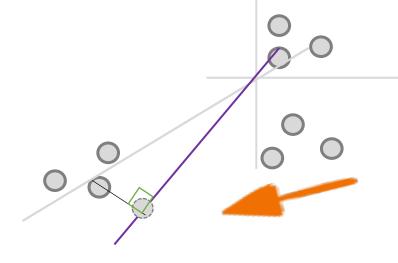
Since PC3 accounts for a very small percentage of overall variance, we can remove it. This is how PCA reduces dimensionality





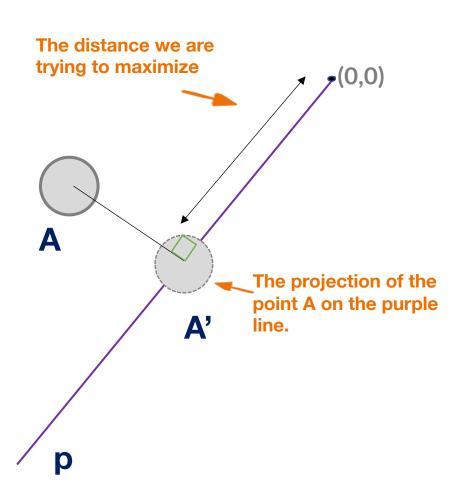


The degree to which a base aligns with the variance represents the amount of information separations along that basis can convey.

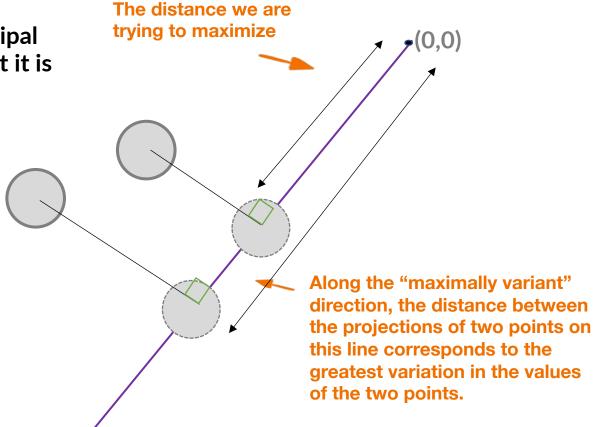


1st base or "Principal Component 1". Line that maximizes sum of distances of projections of points from origin. In essence, maximizes variance of distribution.

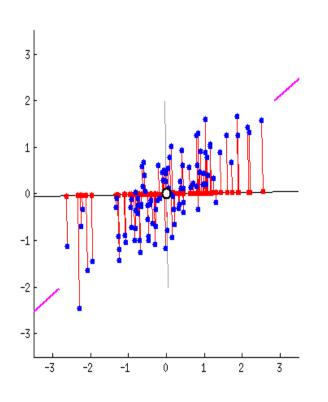
The projection (A') of a point A on a particular line p is the point such that the line AA' is perpendicular to p.



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



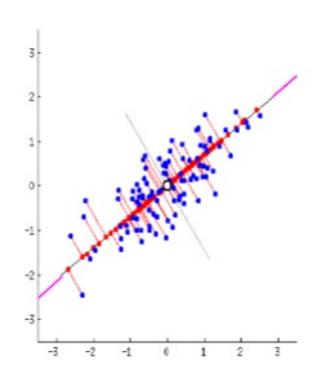
How exactly does maximizing the sum of the distances of these projections from the origin correspond to maximizing the variance along that line?



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



How exactly does maximizing the sum of the distances of these points from the origin correspond to maximizing the variance along that line?



Standardization

**Covariance Matrix Calculation** 

Eigenvector Calculation

Form Principal Components and Build Graph

Individual	Height (cm)	Weight (kg)	Income (\$)	Number of Children
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Compare the data of each of the 4 columns. How do they differ numerically?

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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
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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?



Idea: 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

## **Principal Component Analysis**

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.

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

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## **Principal Component Analysis**

Standardization

**Covariance Matrix Calculation** 

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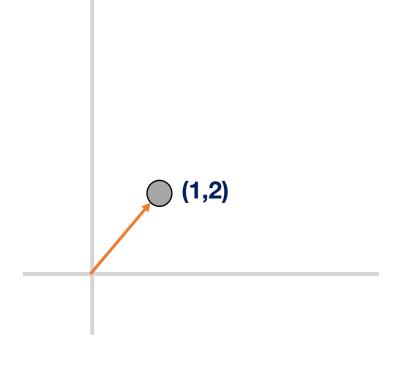
Form Principal Components and Build Graph

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

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

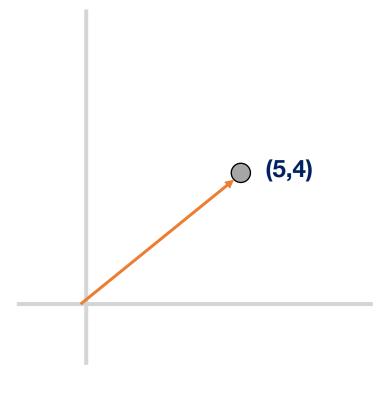


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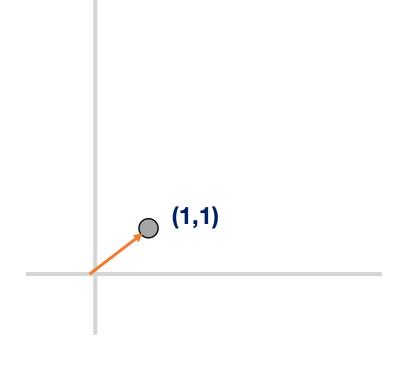
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**Eigenvectors** are characteristic vectors specific to a matrix or transformation.

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

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

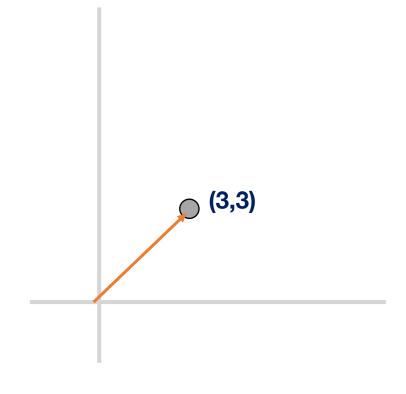


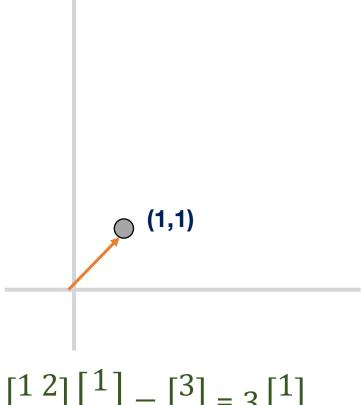
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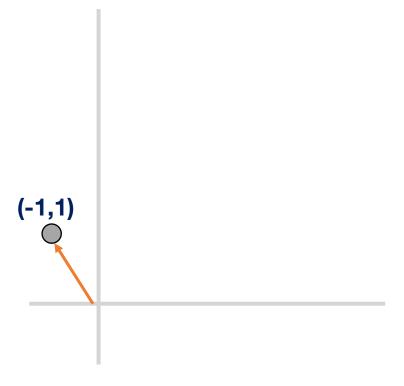
The factor by which an eigenvector is scaled is called its eigenvalue

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

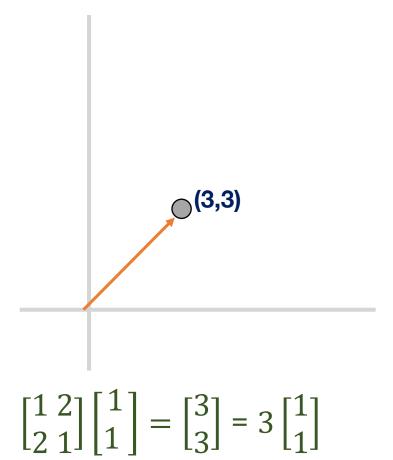


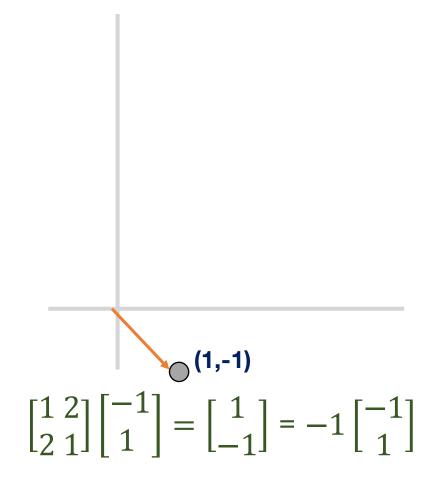


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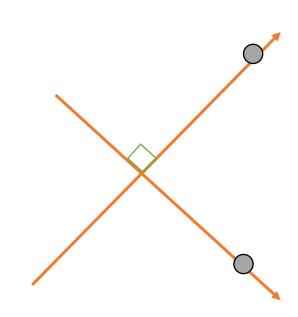


$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} -1 \\ 1 \end{bmatrix} = \begin{bmatrix} 1 \\ -1 \end{bmatrix} = -1 \begin{bmatrix} -1 \\ 1 \end{bmatrix}$$



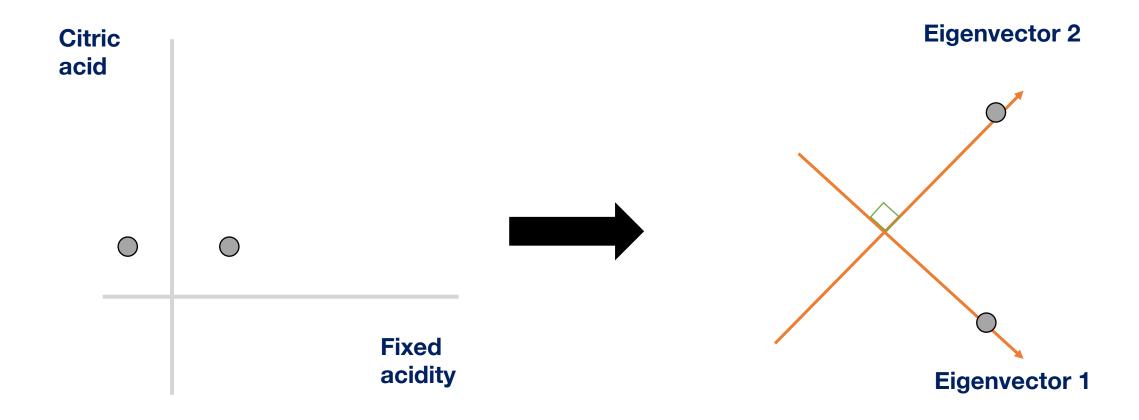


The two eigenvectors are perpendicular to each other!



**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.

## **Principal Component Analysis**

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  $\lambda_1$ ,  $\lambda_2$ ,  $\lambda_3$  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  $\lambda_1$ ,  $\lambda_2$ ,  $\lambda_3$ 

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



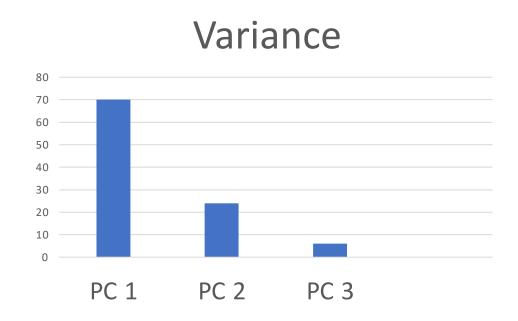
# Form Principal Components and Build Graph

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



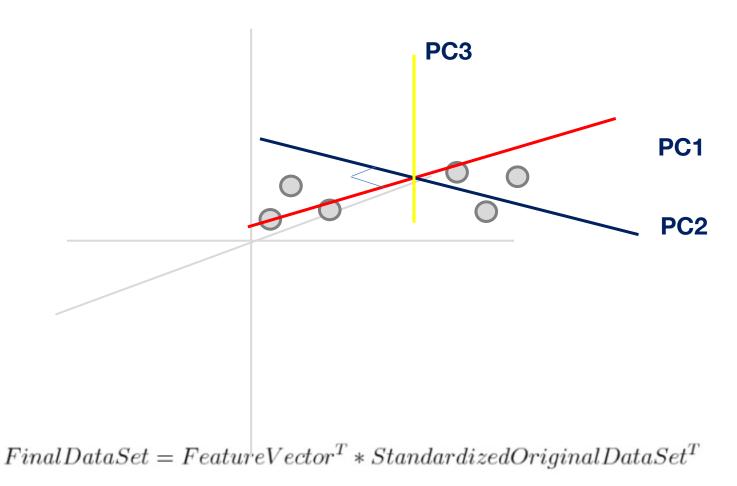
If the percentage of variance of a particular principal component is small enough, discard it. You've now removed a dimension! Form a new matrix which only has the correlation of all the features, eigenvectors/principal components you've selected.

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

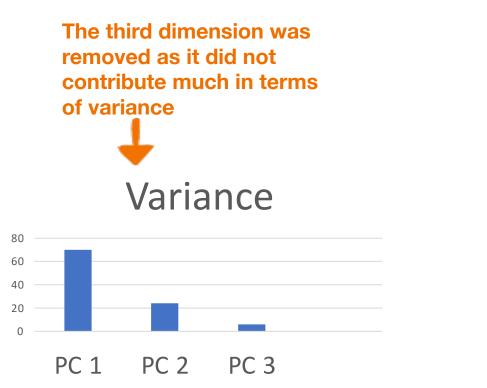


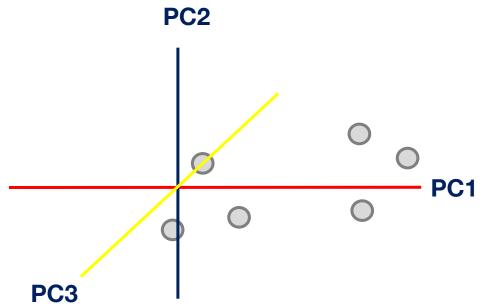
# Now, it's time to reorient the original data along these new axes

## Form Principal Components and Build Graph



# Form Principal Components and Build Graph





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

# A real-world application of PCA

IQ testing!

