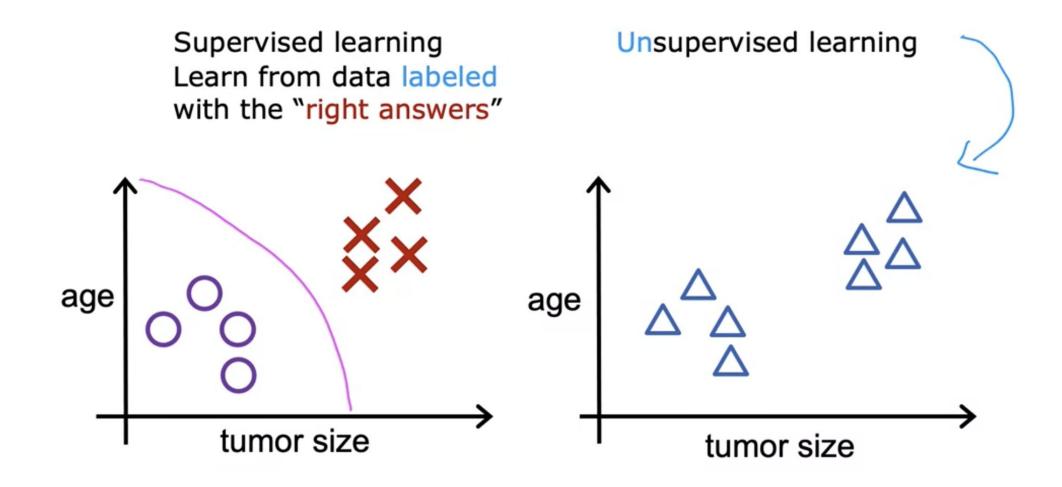




Introducing Unsupervised Learning



Credit: Andrew Ng, Machine 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

View Full Coverage

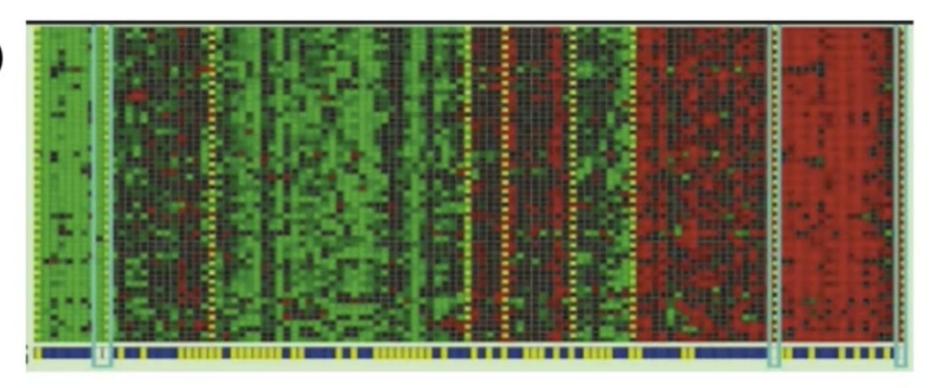
Credit: Andrew Ng, Machine Learning



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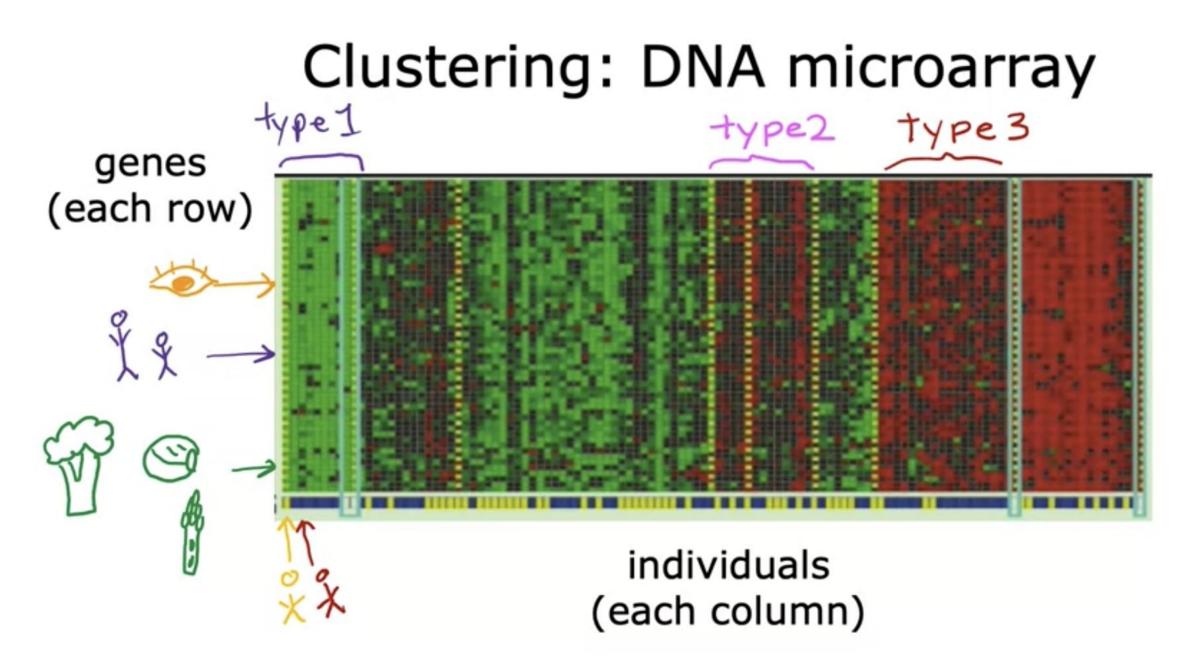
Clustering: DNA microarray

genes (each row)



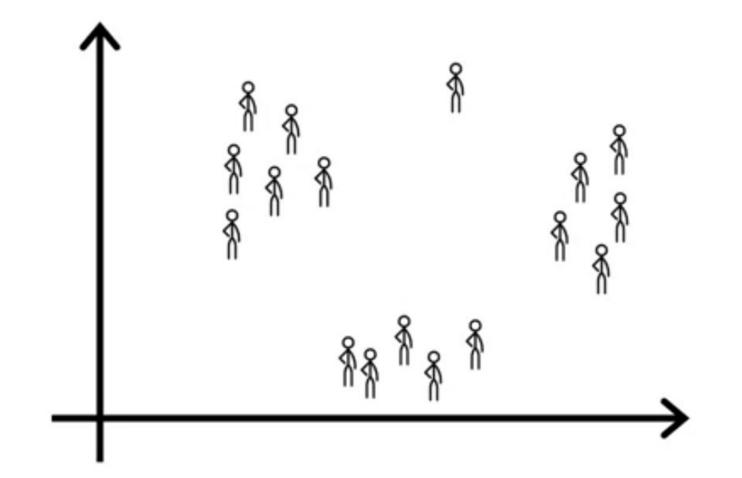
individuals (each column)

Credit: Andrew Ng, Machine Learning



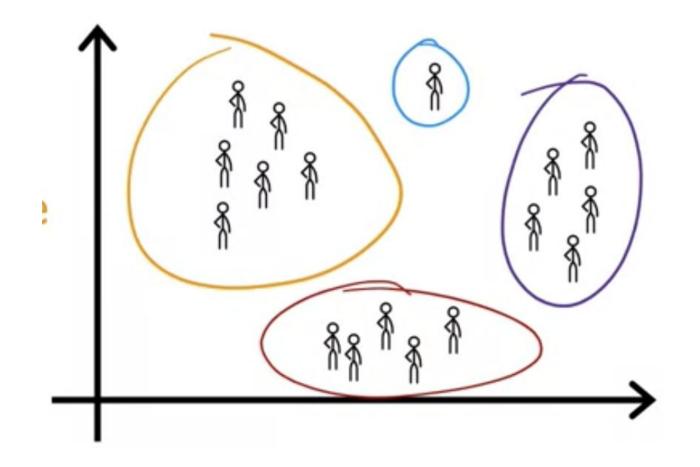
Credit: Andrew Ng, <u>Machine Learning</u>

Clustering: Grouping customers



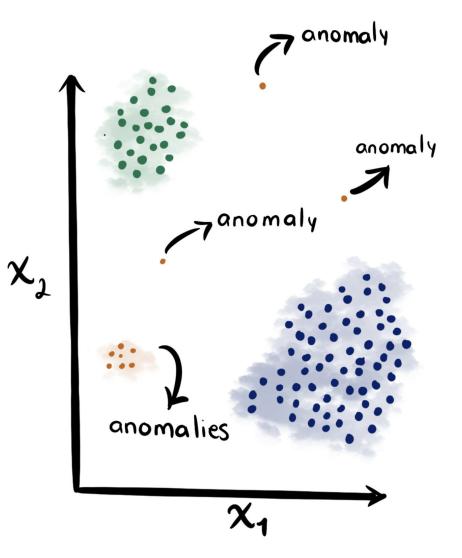
Credit: Andrew Ng, Machine Learning

Grouping Customers

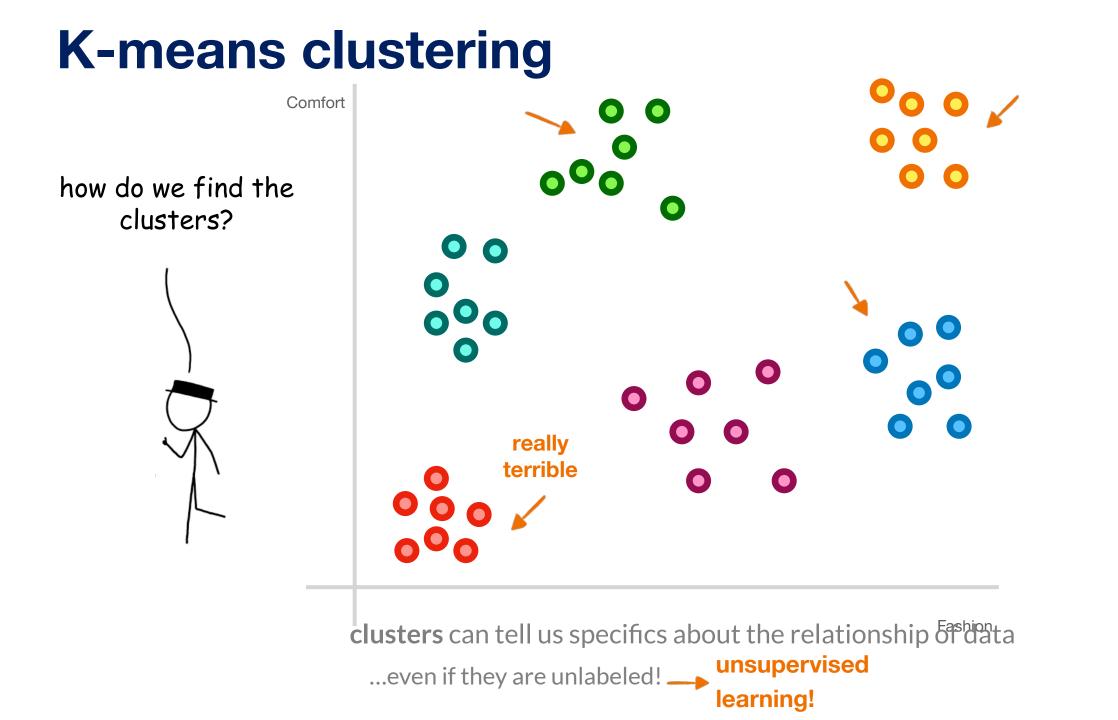


Credit: Andrew Ng, Machine Learning

Anomaly Detection

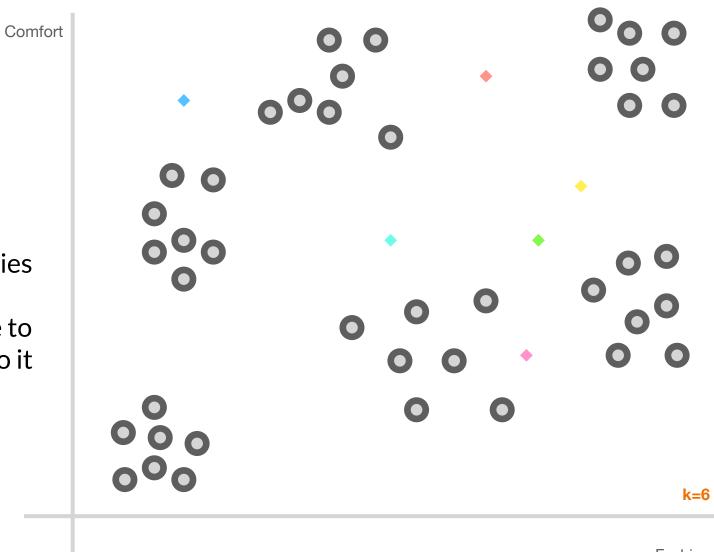


Credit: Anomaly Detection



pick a K-number of clusters randomly pick a series of "centroids" assign each particle to

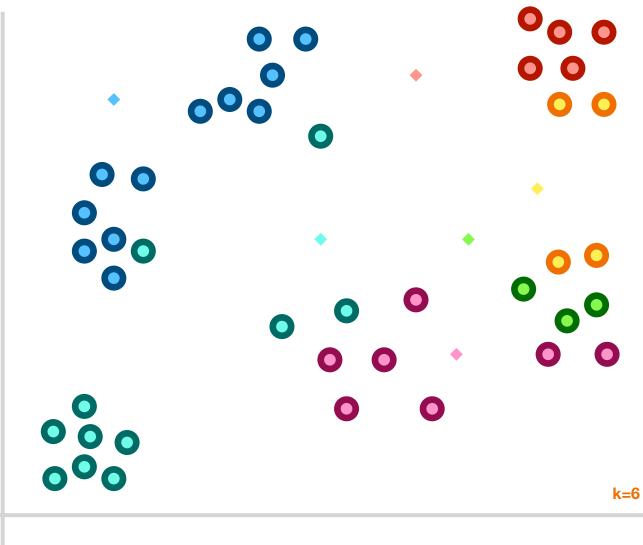
the **centroid** closest to it





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

Comfort

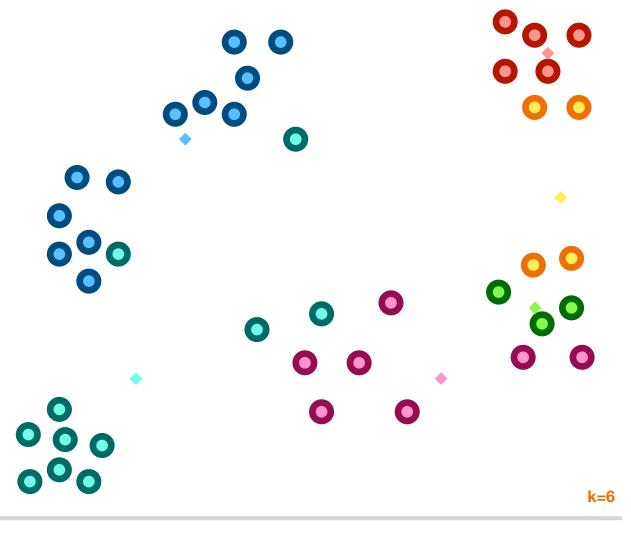


Fashion

pick a K-number of clusters randomly pick a series of "centroids"

Comfort

3. assign each particle to the **centroid** closest to it 4.move the **centroid** to the weighted geometric center of samples assigned to it 5. Repeat 3-4 until centroids stop moving!

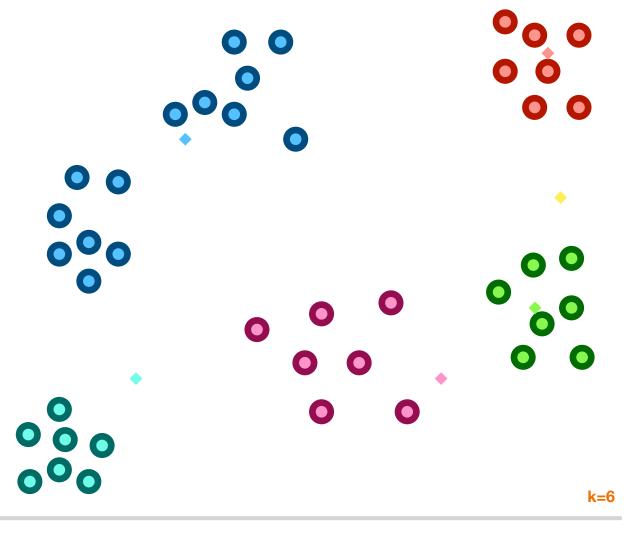


Fashion

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Comfort

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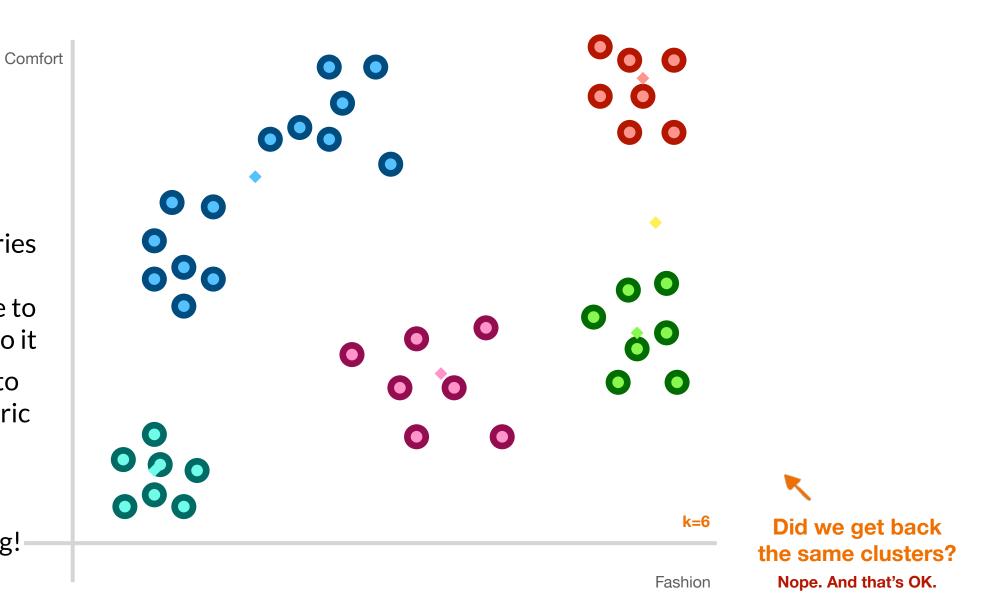


Fashion

pick a K-number of clusters randomly pick a series of "centroids" assign each particle to

the **centroid** closest to it

4.move the **centroid** to the weighted geometric center of samples assigned to it 5. Repeat 3-4 until centroids stop moving!—



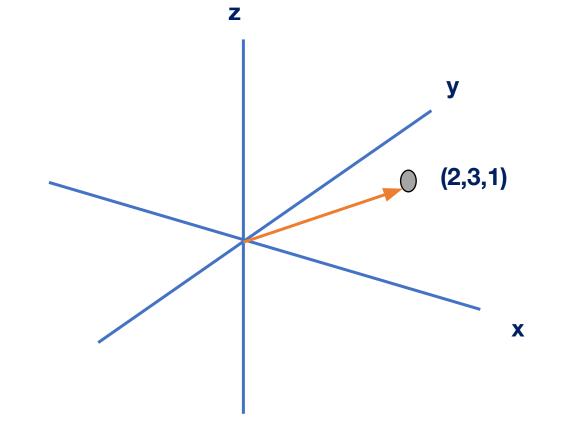
Did we get back the same clusters? Nope. And that's OK.

K-means is an *indeterministic* algorithm—it has built-in randomness

Unsupervised Learning

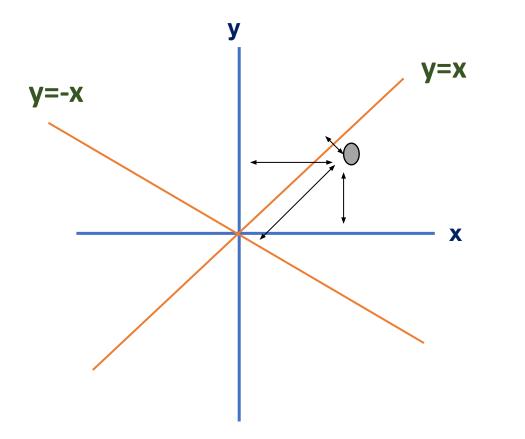
Clustering Dimension reduction

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*.

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

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

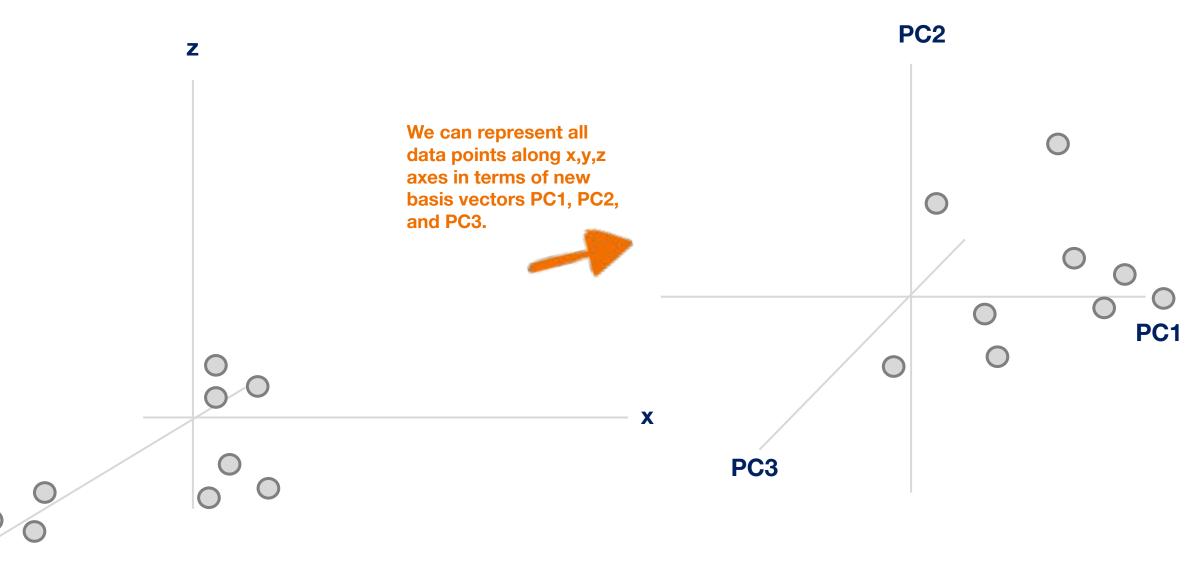
Four dimensions; can't even be graphed!

Motivation for Dimension Reduction

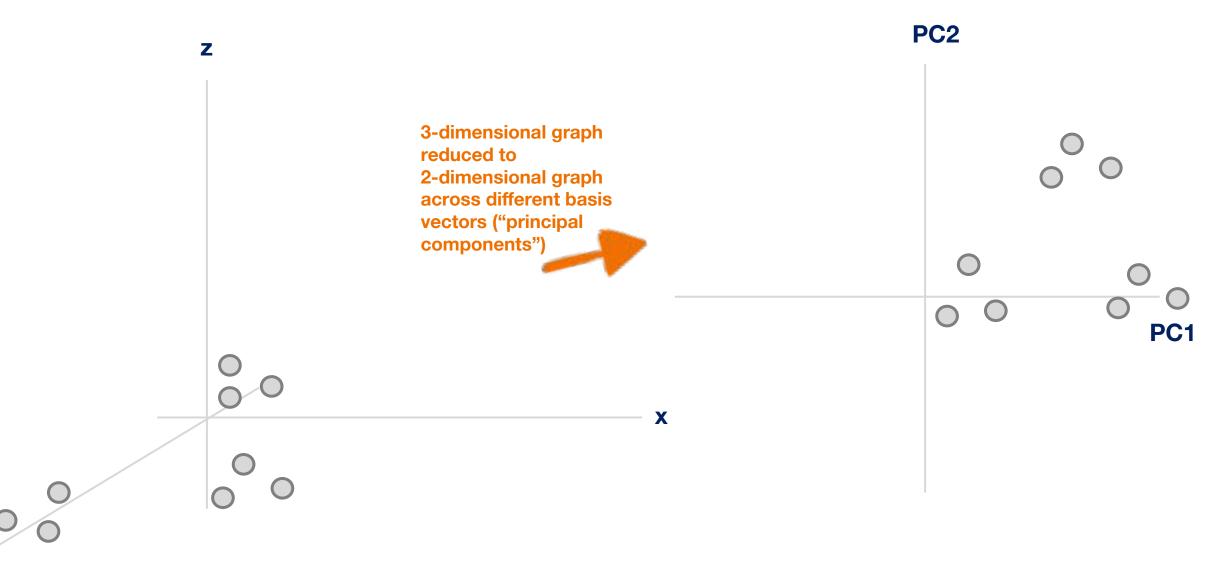
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So, how do we reduce dimensionality without significant loss of information?





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

Principal Components

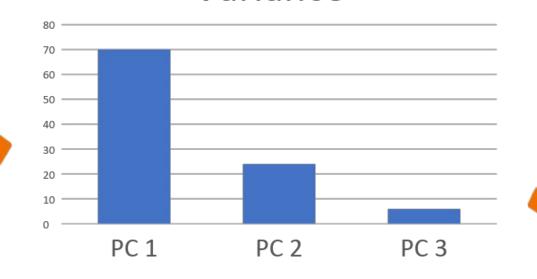
Think of these as new axes that we are orienting our data across.

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, the so that translation across each component indicates different information. **So, they represent directions of maximal variance.**

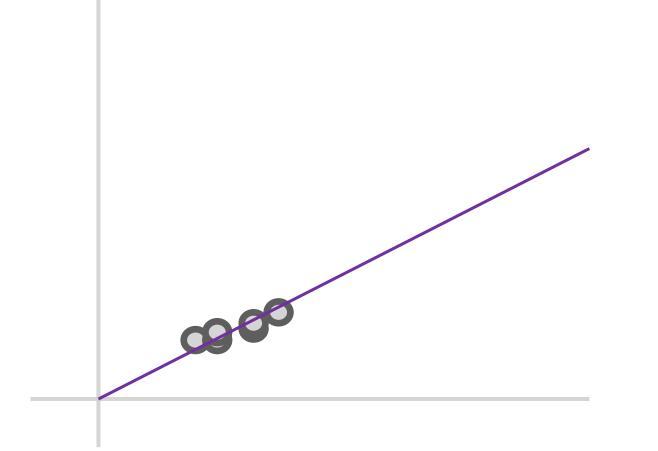
This allows differences between data points to become more prominent **Variance**

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

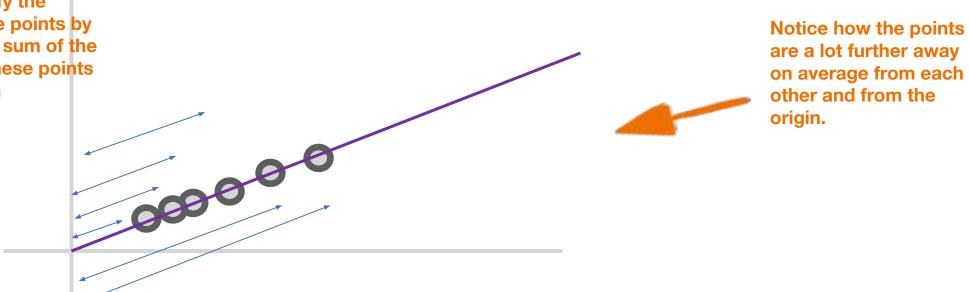


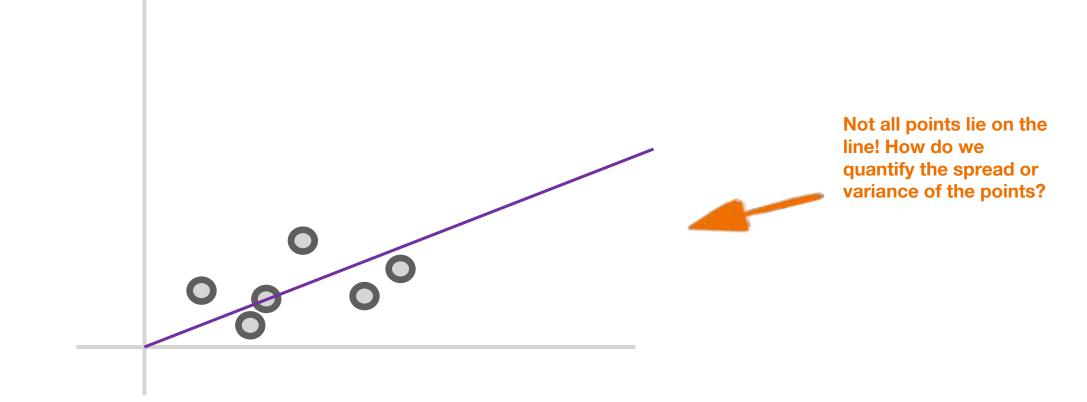
features to remove when reducing the dimensionality of the data? Since PC3 accounts for a very small percentage of overall variance, we can remove it. This is how PCA reduces dimensionality

How do we decide what



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

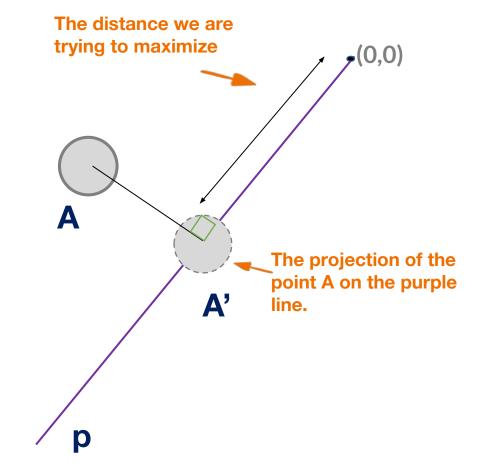




The degree to which a base aligns with the variance represents the amount of information separations along that basis can convey.

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.



The distance we are

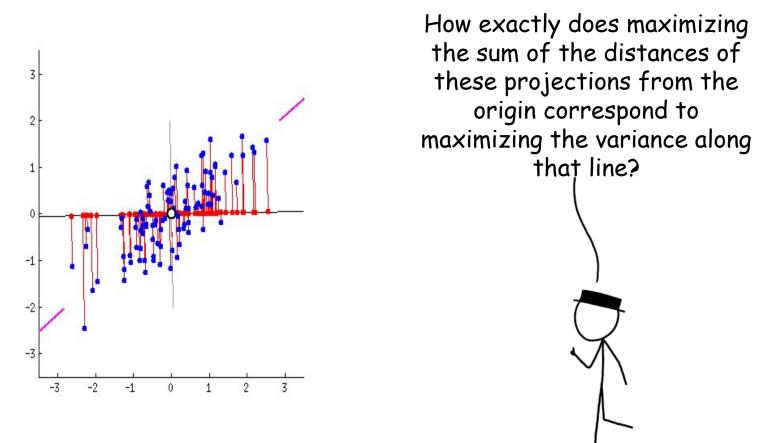
trying to maximize

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

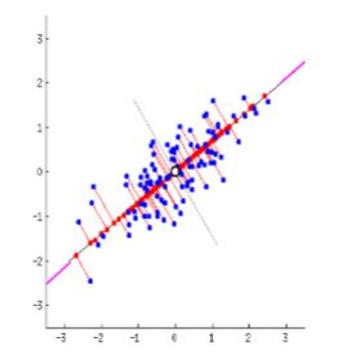
> Along the "maximally variant" direction, the distance between the projections of two points on this line corresponds to the greatest variation in the values of the two points.

(0,0)

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



Built using https://gist.github.com/anonymous/7d888663c6ec679ea6542871 5b99bfdd



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
Person A	165	65	60,000	2
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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.

$$\left[\begin{array}{ccc} Cov(x,x) & Cov(x,y) & Cov(x,z) \\ Cov(y,x) & Cov(y,y) & Cov(y,z) \\ Cov(z,x) & Cov(z,y) & Cov(z,z) \end{array}\right]$$

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

Principal Component Analysis

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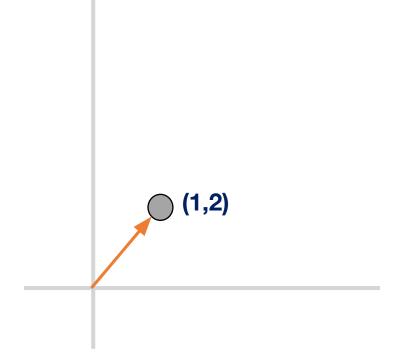
Form Principal Components and Build Graph

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

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

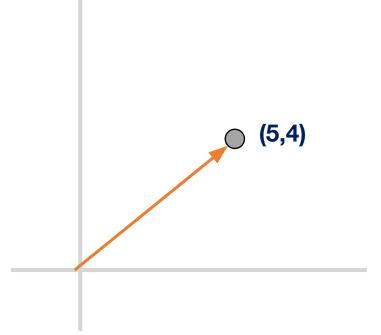


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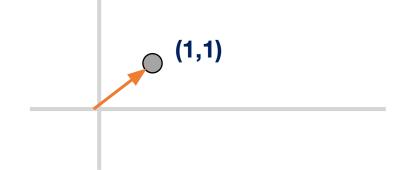
 $\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = \begin{bmatrix} 5 \\ 4 \end{bmatrix}$



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**.

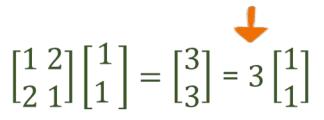
$$\begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \end{bmatrix} = \begin{bmatrix} 3 \\ 3 \end{bmatrix} = 3 \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

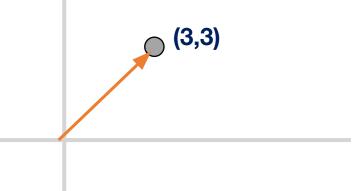


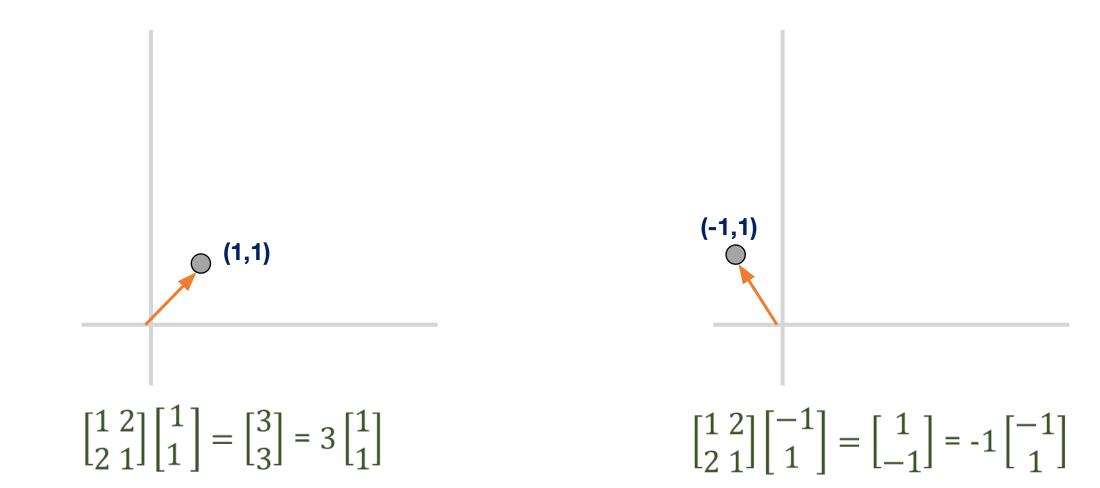
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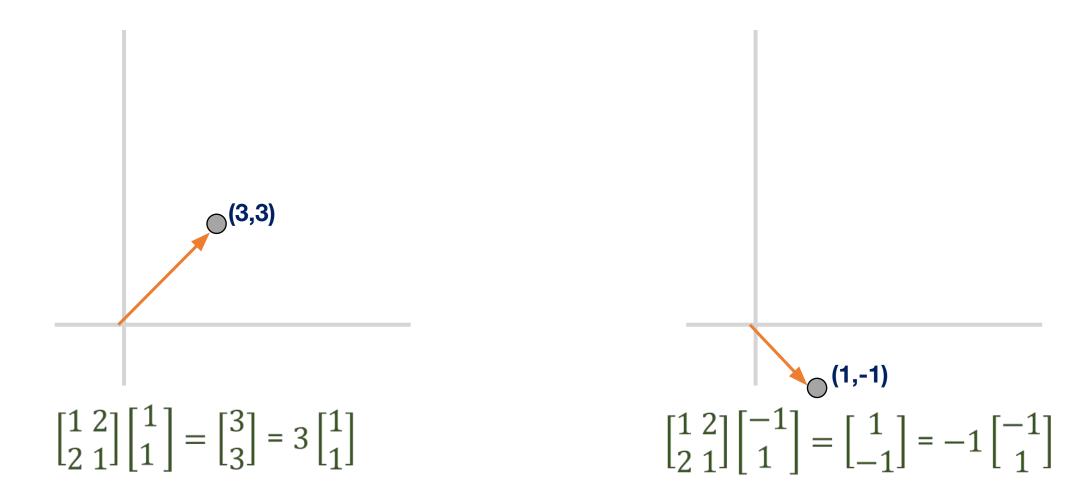
Graphically speaking, when you multiply a matrix with its specific eigenvectors, the eigenvectors **don't get rotated**, **only scaled**.

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







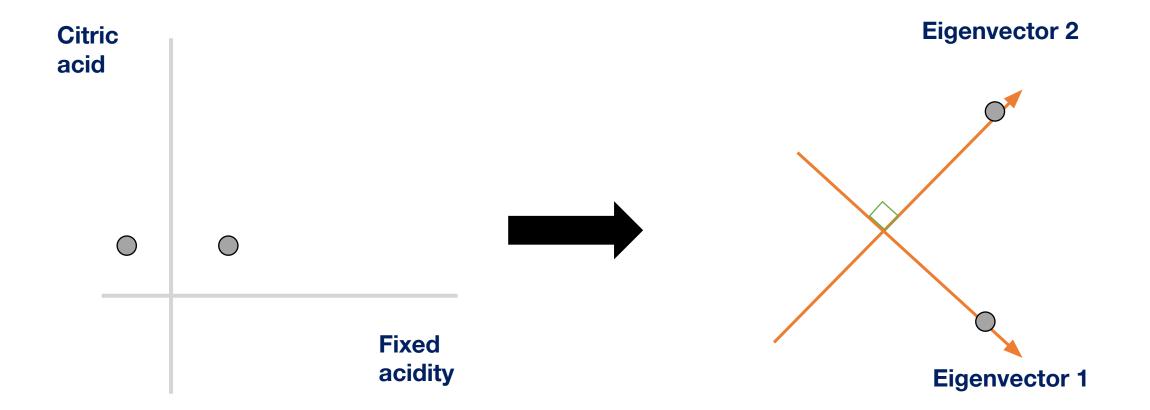


The two eigenvectors are perpendicular to each other!

each

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**.

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

 $Av = \lambda v$

v is the eigenvector and lambda is the eigenvalue

 $Av = \lambda v$

$$\begin{aligned} Av - \lambda v &= 0\\ (A - \lambda)v &= 0\\ |A - \lambda| &= 0 \end{aligned} \qquad A = \begin{bmatrix} Cov(x, x) & Cov(x, y) & Cov(x, z)\\ Cov(y, x) & Cov(y, y) & Cov(y, z)\\ Cov(z, x) & Cov(z, y) & Cov(z, z) \end{bmatrix} \end{aligned}$$

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 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

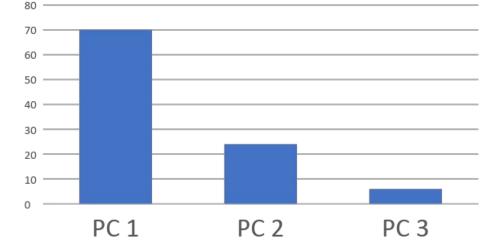
But if the eigenvectors are from the covariance matrix which represents the where will we be removing features?

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**.

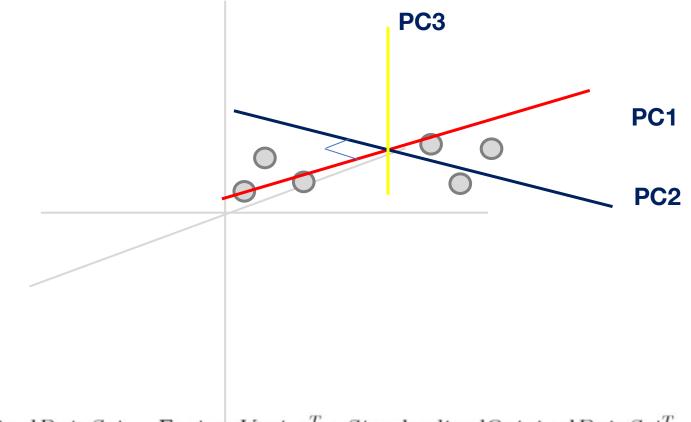






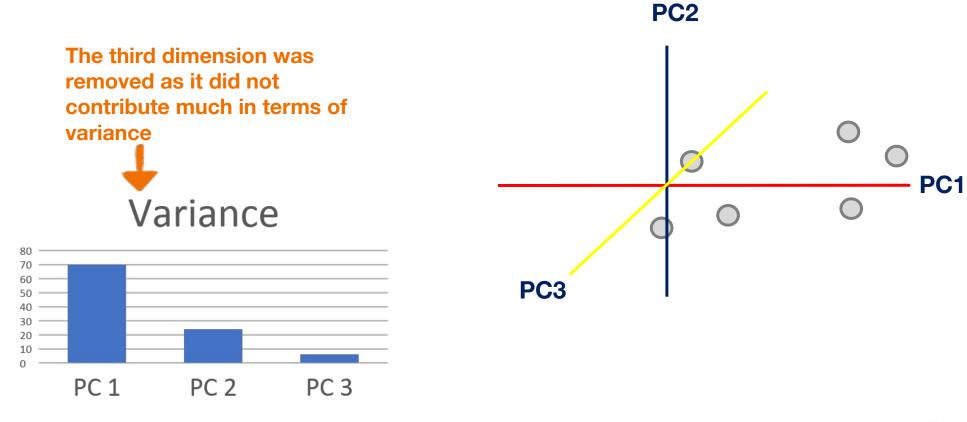
Now, it's time to reorient the original data along these new axes

Form Principal Components and Build Graph



 $FinalDataSet = FeatureVector^{T} * StandardizedOriginalDataSet^{T}$

Form Principal Components and Build Graph



 $FinalDataSet = FeatureVector^{T} * StandardizedOriginalDataSet^{T}$

