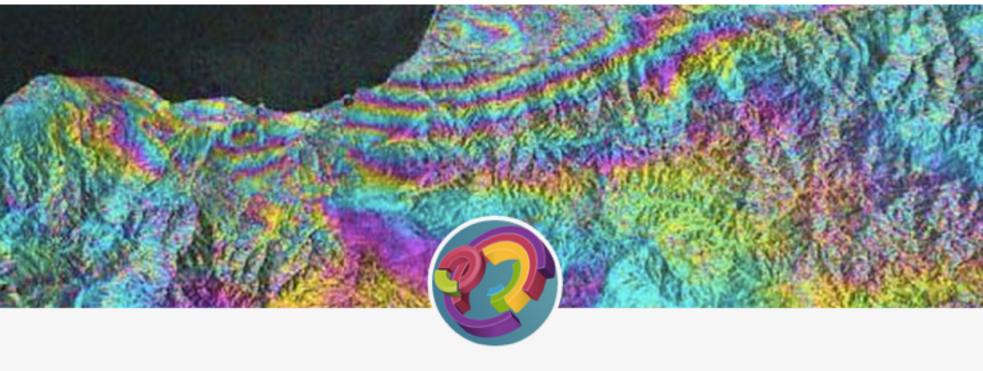
INTRO TO DATA SCIENCE LECTURE 6: DIMENSIONALITY REDUCTION

Francesco Mosconi DAT10 SF // October 22, 2014 INTRO TO DATA SCIENCE, DIMENSIONALITY REDUCTION

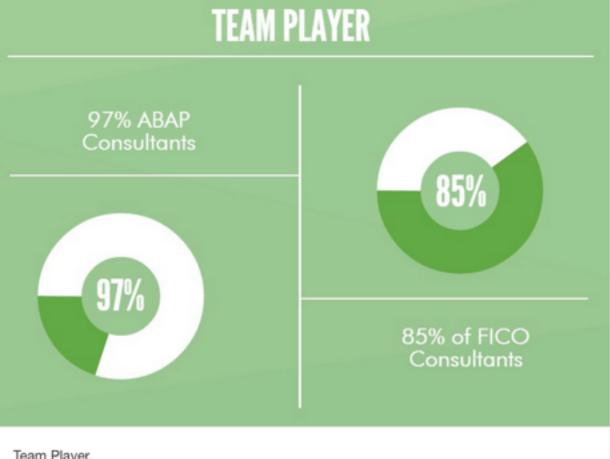
DATA SCIENCE IN THE NEWS



WTF Visualizations

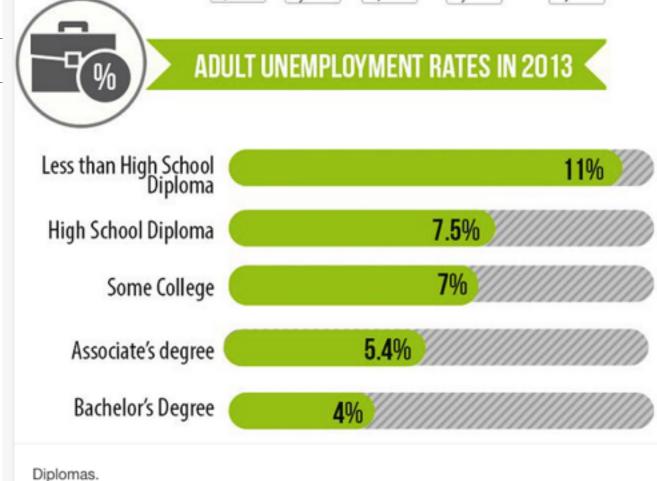
Visualizations that make no sense.

For a discussion of what is wrong with a particular visualization, tweet at us <u>@WTFViz</u>. Check out our friends <u>Thumbs Up Viz</u> and <u>accidental aRt</u>, or <u>submit</u>.

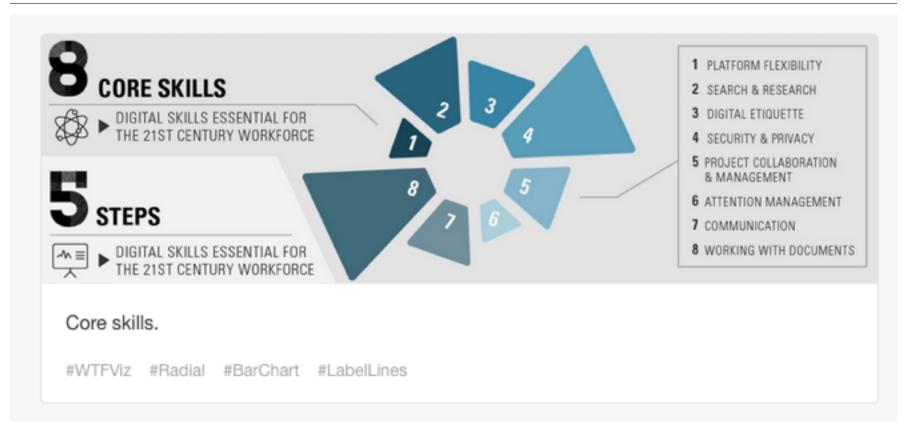


Team Player.

#WTFViz #DonutChart #Percentages



#PartToWhole #Percentages

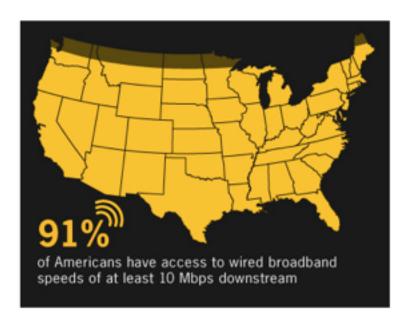


source: http://wtfviz.net/



Inadequate digital skills.

#WTFViz #Clock #PieChart #Percentages



Northern regions.

#WTFViz #Map #Percentages

http://wtfviz.net/

LAST TIME:

- LINEAR REGRESSION (INCL. MULTIPLE REGRESSION)
- POLYNOMIAL REGRESSION
- REGULARIZATION

QUESTIONS?

I. DIMENSIONALITY REDUCTION
II. PRINCIPAL COMPONENTS ANALYSIS
III. SINGULAR VALUE DECOMPOSITION

EXERCISE:
IV. DIMENSIONALITY REDUCTION USING SCIKITLEARN

I. DIMENSIONALITY REDUCTION

A: A set of techniques for reducing the size (in terms of features, records, and/or bytes) of the dataset under examination.

A: A set of techniques for reducing the size (in terms of features, records, and/or bytes) of the dataset under examination.

In general, the idea is to regard the dataset as a matrix and to decompose the matrix into simpler, meaningful pieces.

A: A set of techniques for reducing the size (in terms of features, records, and/or bytes) of the dataset under examination.

In general, the idea is to regard the dataset as a matrix and to decompose the matrix into simpler, meaningful pieces.

Dimensionality reduction is frequently performed as a preprocessing step before another learning algorithm is applied.

	Continuous	Categorical	•
Supervised	???	???	
Unsupervised	???	???	

	Continuous	Categorical
Supervised	regression	classification
Unsupervised	dimension reduction	clustering

Q: What are the motivations for dimensionality reduction?

Q: What are the motivations for dimensionality reduction?

The number of features in our dataset can be difficult to manage, or even misleading (e.g., if the relationships are actually simpler than they appear).

For example, suppose we have a dataset with some features that are related to each other.

For example, suppose we have a dataset with some features that are related to each other.

Ideally, we would like to eliminate this redundancy and consolidate the number of variables we're looking at.

To say this more intuitively, we want to go from a more complex representation of our data to a less complex one (while retaining as much of the signal in our data as possible).

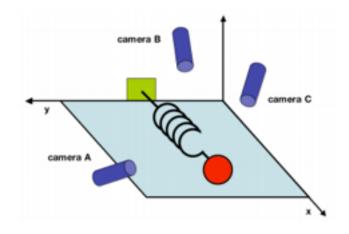
We can do this by looking at our data "from another angle".

In doing this, we tease out the "principal components" of our data.

For example, suppose we have a dataset with some features that are related to each other.

Ideally, we would like to eliminate this redundancy and consolidate the number of variables we're looking at.

If these relationships are *linear*, then we can use well-established techniques like PCA/SVD.



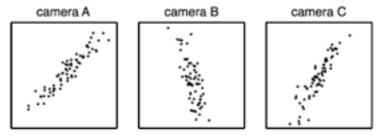


FIG. 1 A toy example. The position of a ball attached to an oscillating spring is recorded using three cameras A, B and C. The position of the ball tracked by each camera is depicted in each panel below.

The complexity that comes with a large number of features is due in part to the curse of dimensionality.

The complexity that comes with a large number of features is due in part to the curse of dimensionality.

Namely, the sample size needed to accurately estimate a random variable taking values in a d-dimensional feature space grows exponentially with d (almost).

Another way of characterizing this is to say that high-dimensional spaces are inherently sparse.

Another way of characterizing this is to say that high-dimensional spaces are inherently sparse.

ex: A high-dimensional orange contains most of its volume in the rind!

ex: A high-dimensional hypercube contains most of its volume in the corners!

In either case, most of the points in the space are "far" from the center.

In either case, most of the points in the space are "far" from the center.

This illustrates the fact that local methods will break down in these circumstances (eg, in order to collect enough neighbors for a given point, you need to expand the radius of the neighborhood so far that locality is not preserved). In either case, most of the points in the space are "far" from the center.

This illustrates the fact that local methods will break down in these circumstances (eg, in order to collect enough neighbors for a given point, you need to expand the radius of the neighborhood so far that locality is not preserved).

The bottom line is that high-dimensional spaces can be problematic.

We'd like to analyze the data using the most meaningful basis (or coordinates) possible.

We'd like to analyze the data using the most meaningful basis (or coordinates) possible.

More precisely: given an $n \times d$ matrix A (encoding n observations of a d-dimensional random variable), we want to find a k-dimensional representation of A (k < d) that captures the information in the original data, according to some criterion.

- reduce computational expense
- reduce susceptibility to overfitting
- reduce noise in the dataset
- enhance our intuition

Q: How is dimensionality reduction performed?

Q: How is dimensionality reduction performed?

A: There are two approaches: feature selection and feature extraction.

Q: How is dimensionality reduction performed?

A: There are two approaches: feature selection and feature extraction.

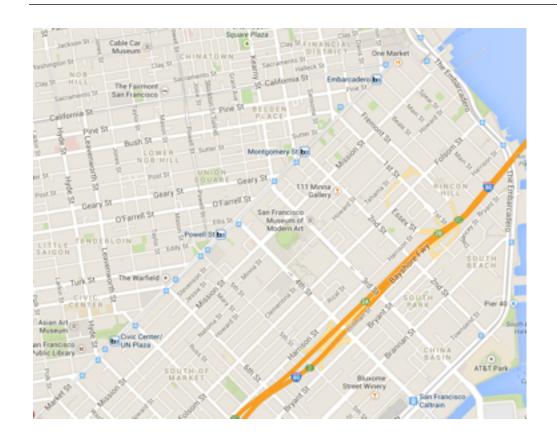
feature selection – selecting a subset of features using an external criterion (filter) or the learning algo accuracy itself (wrapper)

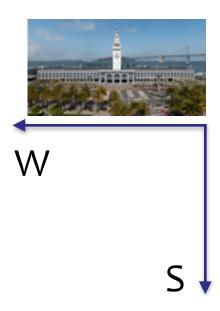
feature extraction – mapping the features to a lower dimensional space

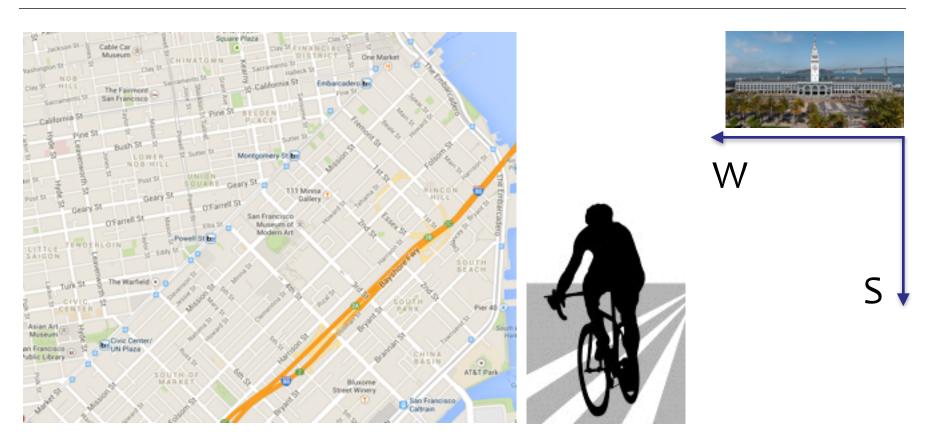
Feature selection is important, but typically when people say dimensionality reduction, they are referring to feature extraction.

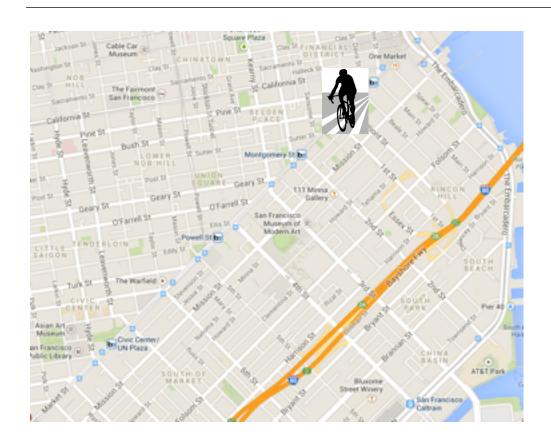
Feature selection is important, but typically when people say dimensionality reduction, they are referring to feature extraction.

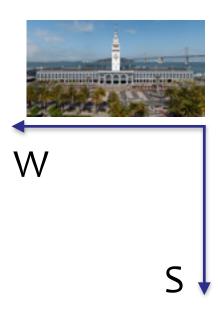
The goal of feature extraction is to create a new set of coordinates that simplify the representation of the data.

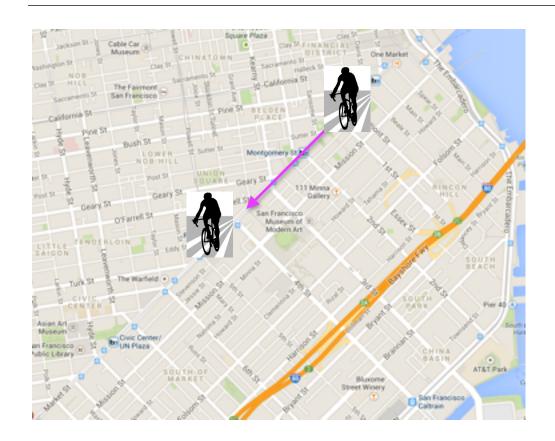


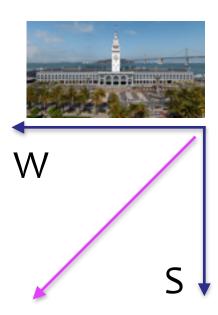


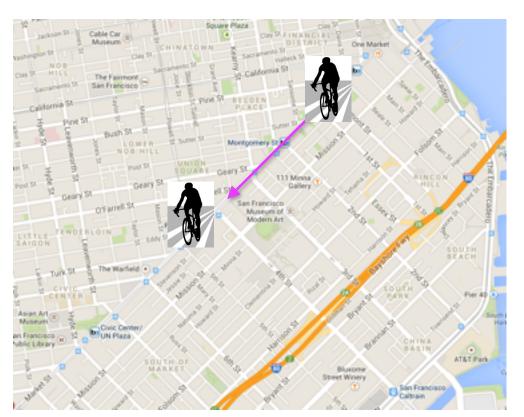


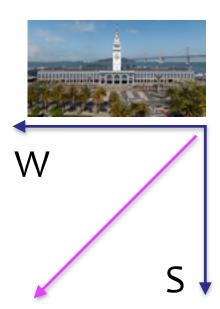




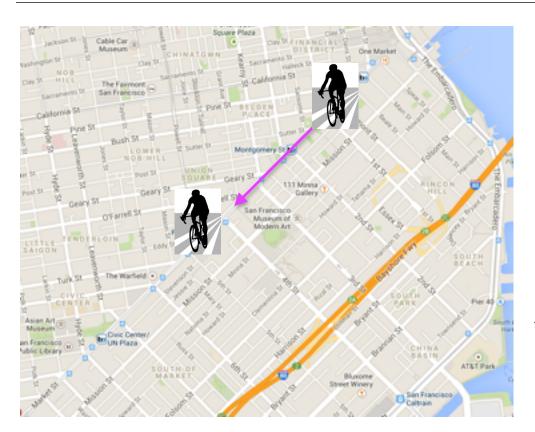


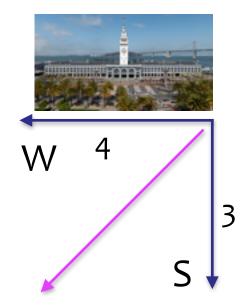




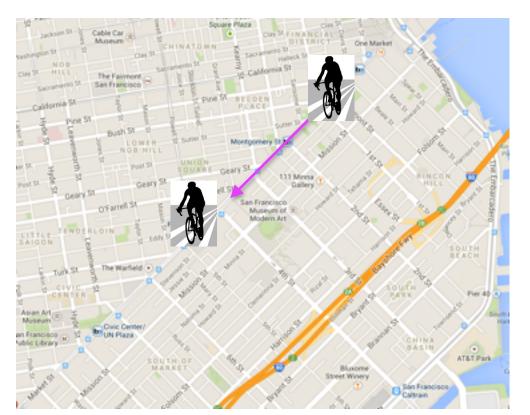


How many dimensions do we need to specify the position of this bike?

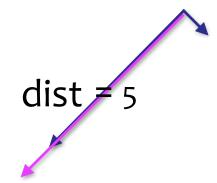




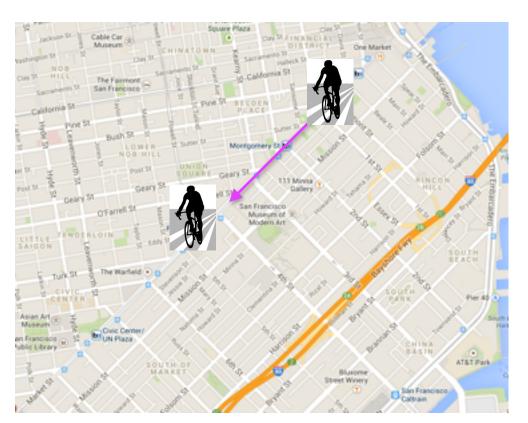
Yep, two. But could we represent the biker's position with fewer dimensions? How?

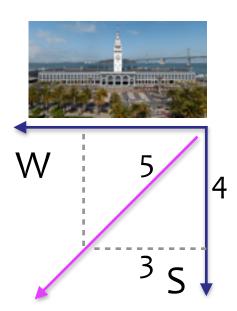






What if we just used distance down Market St.?





Of course, we can always map back to the original coordinate system!

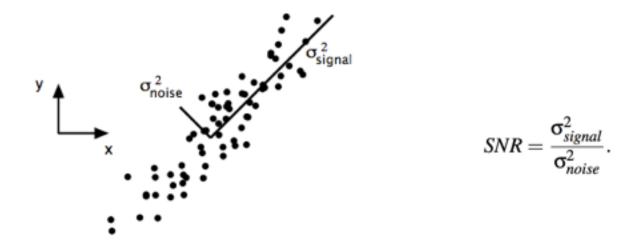
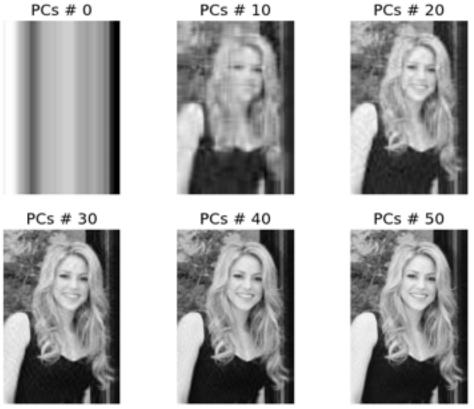


FIG. 2 Simulated data of (x,y) for camera A. The signal and noise variances σ_{signal}^2 and σ_{noise}^2 are graphically represented by the two lines subtending the cloud of data. Note that the largest direction of variance does not lie along the basis of the recording (x_A, y_A) but rather along the best-fit line.

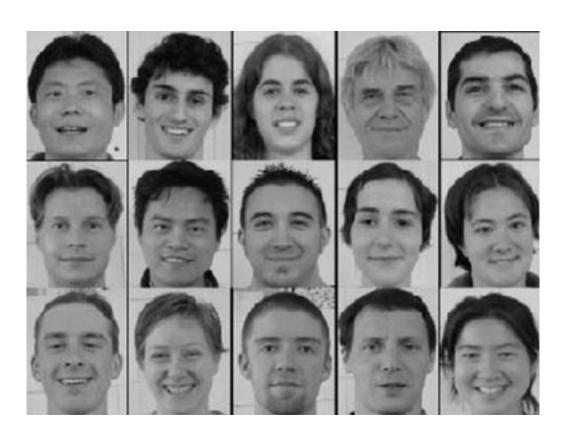
Q: What are some applications of dimensionality reduction?

Q: What are some applications of dimensionality reduction?

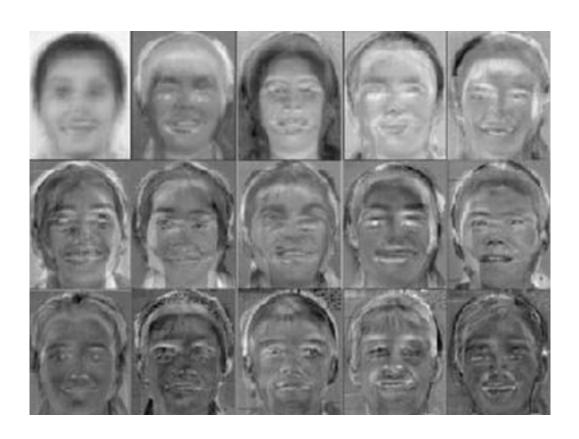
- topic models (document clustering)
- image recognition/computer vision
- bioinformatics (microarray analysis)
- speech recognition
- astronomy (spectral data analysis)
- recommender systems



source: http://glowingpython.blogspot.it/2011/07/pca-and-image-compression-with-numpy.html



DIMENSIONALITY REDUCTION



II. PRINCIPAL COMPONENT ANALYSIS

Principal component analysis is a dimension reduction technique that can be used on a matrix of any dimensions.

Principal component analysis is a dimension reduction technique that can be used on a matrix of any dimensions.

This procedure produces a new basis (a new coordinate system), each of whose components retain as much variance from the original data as possible.

Principal component analysis is a dimension reduction technique that can be used on a matrix of any dimensions.

This procedure produces a new basis, each of whose components retain as much variance from the original data as possible.

The PCA of a matrix A boils down to the eigenvalue decomposition of the covariance matrix of A.

The covariance matrix C of a matrix A is always square:

$$C = \begin{bmatrix} E[(X_1 - \mu_1)(X_1 - \mu_1)] & E[(X_1 - \mu_1)(X_2 - \mu_2)] & \cdots & E[(X_1 - \mu_1)(X_n - \mu_n)] \\ E[(X_2 - \mu_2)(X_1 - \mu_1)] & E[(X_2 - \mu_2)(X_2 - \mu_2)] & \cdots & E[(X_2 - \mu_2)(X_n - \mu_n)] \\ \vdots & \vdots & \ddots & \vdots \\ E[(X_n - \mu_n)(X_1 - \mu_1)] & E[(X_n - \mu_n)(X_2 - \mu_2)] & \cdots & E[(X_n - \mu_n)(X_n - \mu_n)] \end{bmatrix}.$$

off-diagonal elements C_{ij} give the covariance between X_i , X_j ($i \neq j$) diagonal elements C_{ii} give the variance of X_i

Wait a minute, what's a covariance matrix?

$$\mathbf{c} = \begin{bmatrix} \mathbf{E}[(X_1 - \mu_1)(X_1 - \mu_1)] & \mathbf{E}[(X_1 - \mu_1)(X_2 - \mu_2)] & \cdots & \mathbf{E}[(X_1 - \mu_1)(X_n - \mu_n)] \\ \mathbf{E}[(X_2 - \mu_2)(X_1 - \mu_1)] & \mathbf{E}[(X_2 - \mu_2)(X_2 - \mu_2)] & \cdots & \mathbf{E}[(X_2 - \mu_2)(X_n - \mu_n)] \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{E}[(X_n - \mu_n)(X_1 - \mu_1)] & \mathbf{E}[(X_n - \mu_n)(X_2 - \mu_2)] & \cdots & \mathbf{E}[(X_n - \mu_n)(X_n - \mu_n)] \end{bmatrix}.$$

For that matter, what is covariance?

Remember variance?

Remember variance?

$$s^{2} = \frac{\sum_{i=1}^{n} (X_{i} - X)^{2}}{(n-1)}$$

Variance is the average distance from the mean of a data set to a point in that data set.

In other words, it is a measure of the spread of the data.

Recall that standard deviation is the square root of variance.

Standard deviation and variance only operate on 1 dimension, so that you could only calculate the standard deviation for each dimension of the data set *independently* of the other dimensions. However, it is useful to have a similar measure to find out how much the dimensions vary from the mean with respect to each other.

This is called covariance.

Variance:

$$s^{2} = \frac{\sum_{i=1}^{n} (X_{i} - \bar{X})^{2}}{(n-1)} \qquad var(X) = \frac{\sum_{i=1}^{n} (X_{i} - \bar{X})(X_{i} - \bar{X})}{(n-1)}$$

Covariance:

$$cov(X,Y) = \frac{\sum_{i=1}^{n} (X_i - \bar{X})(Y_i - \bar{Y})}{(n-1)}$$

Covariance is always measured between two dimensions. If you calculate the covariance between a dimension and itself, you get the variance.

The covariance matrix C of a matrix A is always square:

$$C = \begin{pmatrix} cov(x,x) & cov(x,y) & cov(x,z) \\ cov(y,x) & cov(y,y) & cov(y,z) \\ cov(z,x) & cov(z,y) & cov(z,z) \end{pmatrix}$$

off-diagonal elements C_{ij} give the covariance between X_i , X_j ($i \neq j$) diagonal elements C_{ii} give the variance of X_i

The covariance matrix C of a matrix A is always square:

$$C = \begin{bmatrix} E[(X_1 - \mu_1)(X_1 - \mu_1)] & E[(X_1 - \mu_1)(X_2 - \mu_2)] & \cdots & E[(X_1 - \mu_1)(X_n - \mu_n)] \\ E[(X_2 - \mu_2)(X_1 - \mu_1)] & E[(X_2 - \mu_2)(X_2 - \mu_2)] & \cdots & E[(X_2 - \mu_2)(X_n - \mu_n)] \\ \vdots & \vdots & \ddots & \vdots \\ E[(X_n - \mu_n)(X_1 - \mu_1)] & E[(X_n - \mu_n)(X_2 - \mu_2)] & \cdots & E[(X_n - \mu_n)(X_n - \mu_n)] \end{bmatrix}.$$

off-diagonal elements C_{ij} give the covariance between X_i , X_j ($i \neq j$) diagonal elements C_{ii} give the variance of X_i

$$A = Q\Lambda Q^{-1}$$

$$A = Q\Lambda Q^{-1}$$

The columns of Q are the eigenvectors of A, and the values in Λ are the associated eigenvalues of A.

$$A = Q\Lambda Q^{-1}$$

The columns of Q are the eigenvectors of A, and the values in Λ are the associated eigenvalues of A.

For an eigenvector v of A and its eigenvalue λ , we have the important relation:

$$AV = \lambda V$$

$$A = Q\Lambda Q^{-1}$$

The columns of Q are the eigenvectors of A, and the are the associated eigenvalues of A.

NOTE

This relationship defines what it means to be an eigenvector of Δ

For an eigenvector V of A and its eigenvalue λ , we have the important relation:

$$Av = \lambda v$$

The eigenvectors form a basis of the vector space on which *A* acts (e.g., they are orthogonal).

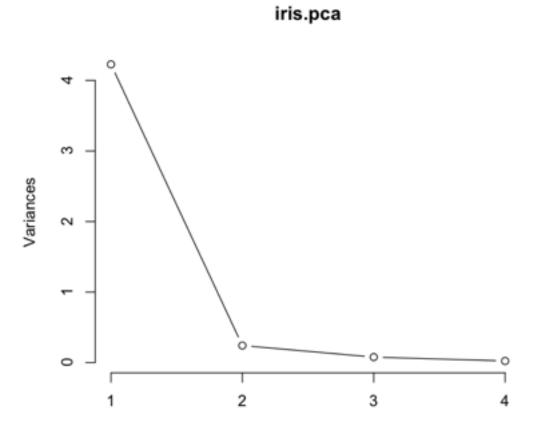
The eigenvectors form a basis of the vector space on which *A* acts (e.g., they are orthogonal).

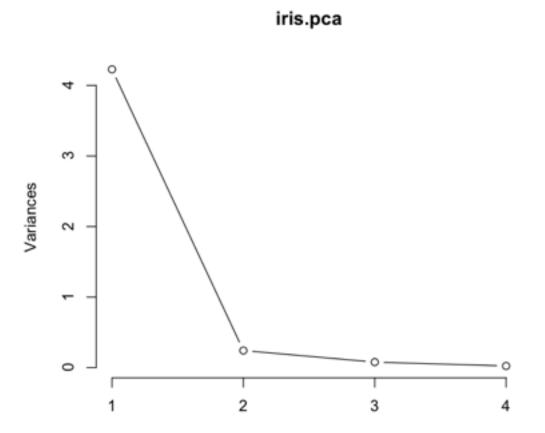
Furthermore the basis elements are ordered by their eigenvalues (from largest to smallest), and these eigenvalues represent the amount of variance explained by each basis element.

The eigenvectors form a basis of the vector space on which *A* acts (eg, they are orthogonal).

Furthermore the basis elements are ordered by their eigenvalues (from largest to smallest), and these eigenvalues represent the amount of variance explained by each basis element.

This can be visualized in a scree plot, which shows the amount of variance explained by each basis vector.





NOTE

Looking at this plot also gives you an idea of how many principal components to keep.

Apply the *elbow test*: keep only those pc's that appear to the left of the elbow in the graph.

- 1. Linearity The change in basis is a <u>linear</u> projection
- 2. Large variances have important structure e.g. large signal-to-noise ratio. In other words, we assume that principal components with larger associated variances are signal, while those with lower variances represent noise. NOTE: this is a strong (and not always correct) assumption!
- 3. The principal components are orthogonal A simplification that makes PCA soluble with linear algebra matrix decomposition techniques

III. SINGULAR VALUE DECOMPOSITION

Notice: Lots of math / linear algebra notation ahead!

It's okay if it does not all immediately make sense.

Take a deep breath...



Notice: Lots of math / linear algebra notation ahead!

It's okay if it does not all immediately make sense.

Take a deep breath...

That's better! Okay, then...



KEEP CALM AND TAKE A DEEP BREATH

The singular value decomposition of *A* is given by:

$$A = U \Sigma V^T$$

The singular value decomposition of *A* is given by:

$$A = U \sum_{(n \times d)} V^{T}$$

The singular value decomposition of *A* is given by:

$$A = U \sum V^{T}$$
(n x d) (n x n) (n x d) (d x d)

st. U, V are orthogonal matrices and Σ is a diagonal matrix.

The singular value decomposition of *A* is given by:

$$A = U \sum_{(n \times d)} V^{T}$$

st. U, V are orthogonal matrices and Σ is a diagonal matrix.

The singular value decomposition of A is given by:

$$A = U \sum V^{T}$$
(n x d) (n x d) (d x d)

The columns of U & V are the (left- and right-) singular vectors of A.

The singular value decomposition of A is given by:

$$A = U \sum_{(n \times d)} V^{T}$$

The columns of U&V are the (left- and right-) singular vectors of A.

These singular vectors provide orthonormal bases for the spaces $K_n \& K_d$ (columns of U & V, respectively).

The singular value decomposition of A is given by:

$$A = U \sum_{(n \times d)} V^{T}$$

The nonzero entries of Σ are the singular values of A. These are real, nonnegative, and rank-ordered (decreasing from left to right).

For a general SVD, the columns of U are the eigenvectors of AA^T , and the columns of V are the eigenvectors of A^TA .

Also, the singular values of A are the square roots of the eigenvalues of AA^{T} and $A^{T}A$.

Q: How do you interpret the SVD?

Q: How do you interpret the SVD?

A: Recall that given a set of n points in d-dimensional space (e.g., a matrix A), we want to find the best k < d dimensional subspace to represent the data.

Q: How do you interpret the SVD?

A: Recall that given a set of n points in d-dimensional space (eg, a matrix A), we want to find the best k < d dimensional subspace to represent the data.

For k = 1, this subspace is a line passing through the origin.

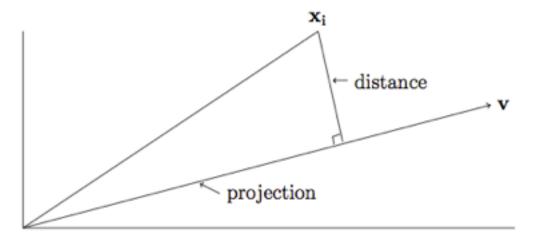


Figure 4.1: The projection of the point $\mathbf{x_i}$ onto the line through the origin in the direction of \mathbf{v}

IV. OTHER METHODS

Whereas PCA and SVD create new coordinates by transform the old coordinates, factor analysis requires new coordinates to be specified externally.

Whereas PCA and SVD create new coordinates by transform the old coordinates, factor analysis requires new coordinates to be specified externally.

These new coordinates are associated with hidden or latent features that we think our data depends on.

Whereas PCA and SVD create new coordinates by transform the old coordinates, factor analysis requires new coordinates to be specified externally.

These new coordinates are associated with hidden or latent features that we think our data depends on.

The old coordinates are then modeled as linear combinations of the latent features. For example, consider a dataset that represents the results of a decathalon (rows = participants, columns = events, entries = times).

For example, consider a dataset that represents the results of a decathalon (rows = participants, columns = events, entries = times).

Though this dataset contains 10 features X_i , we may be interested in modeling these features as functions of *latent* variables such as the speed and strength of the participants:

$$X_i = \lambda_1 f_1 + \lambda_2 f_2 + \varepsilon$$

For example, consider a dataset that represents the results of a decathalon (rows = participants, columns = events, entries = times).

Though this dataset contains 10 features X_i , we may be interested in modeling these features as functions of *latent* variables such as the speed and strength of the participants:

$$X_i = \lambda_1 f_1 + \lambda_2 f_2 + \varepsilon$$

This would allow us to analyze the data in a more fundamental way.

SVD, PCA, and factor analysis are all linear techniques (eg, we use a linear transformation to embed the in a lower-dimensional space).

However, sometimes linear techniques are not sufficient.

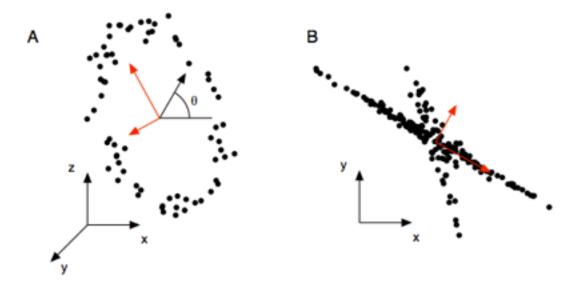


FIG. 6 Example of when PCA fails (red lines). (a) Tracking a person on a ferris wheel (black dots). All dynamics can be described by the phase of the wheel θ , a non-linear combination of the naive basis. (b) In this example data set, non-Gaussian distributed data and non-orthogonal axes causes PCA to fail. The axes with the largest variance do not correspond to the appropriate answer.