INTRO TO DATA SCIENCE DIMENSIONALITY REDUCTION

I. DIMENSIONALITY REDUCTION
II. PRINCIPAL COMPONENTS ANALYSIS
III. BONUS SINGULAR VALUE DECOMPOSITION
IV. BONUS OTHER METHODS

EXERCISE:

IV. DIMENSIONALITY REDUCTION IN SCIKIT-LEARN

| continuous categorical | Supervised |

Unsupervised

Supervised regression classification
Unsupervised dimension reduction

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Dimensionality reduction is frequently performed as a pre-processing step before another learning algorithm is applied.

Q: What are the motivations for dimensionality reduction?

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

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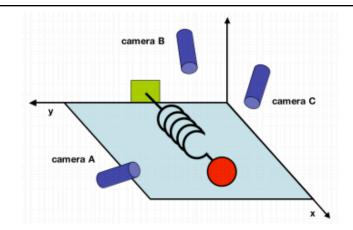
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If these relationships are *linear*, then we can use well-established techniques like PCA/SVD.

EXAMPLE: 1D HARMONIC OSCILLATOR



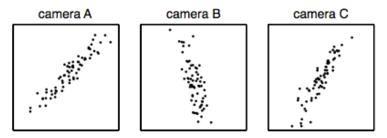


FIG. 1 A toy example. The position of a ball attached to an oscillating spring is recorded using three cameras A, B and C. The position of the ball tracked by each camera is depicted in each panel below.

ASIDE: CURSE OF DIMENSIONALITY

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

(More precisely, the sample size grows exponentially with $l \le d$, the dimension of the manifold *embedded* in the feature space).

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

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

Q: What is the goal of dimensionality reduction?

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

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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*.

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

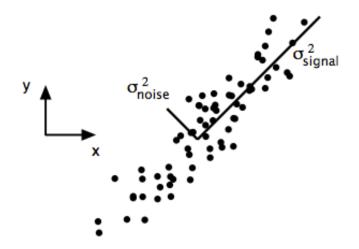


FIG. 2 Simulated data of (x,y) for camera A. The signal and noise variances σ_{signal}^2 and σ_{noise}^2 are graphically represented by the two lines subtending the cloud of data. Note that the largest direction of variance does not lie along the basis of the recording (x_A, y_A) but rather along the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

Q: What are some applications of dimensionality reduction?

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





PCs # 10



PCs # 40



PCs # 20



PCs # 50



II. PRINCIPAL COMPONENT ANALYSIS

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

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

The covariance matrix C of a matrix X 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

ASIDE: EIGENVALUE DECOMPOSITION

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

$$Cv = \lambda v$$

The eigenvectors form a basis of the vector space on which ${\cal C}$ acts (eg, they are orthogonal).

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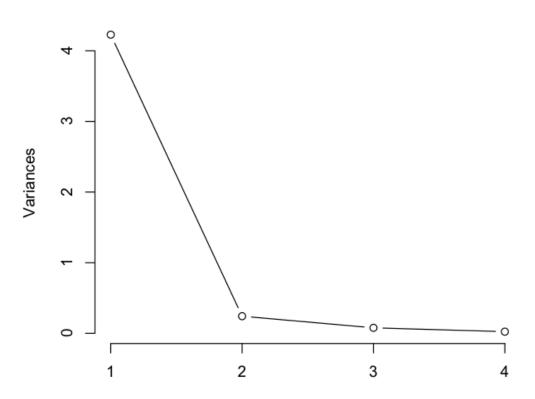
Furthermore the basis elements are ordered by their eigenvalues (from largest to smallest), and these eigenvalues represent the amount of variance explained by each basis element.

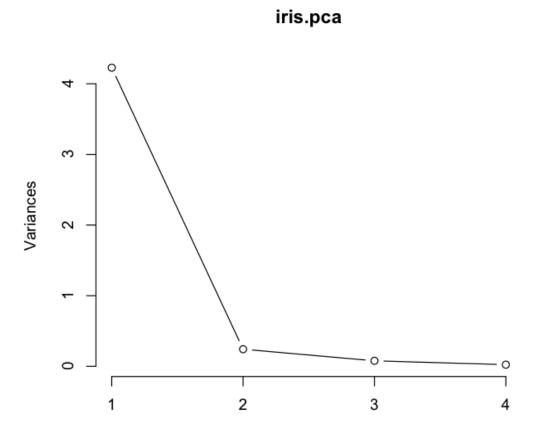
The eigenvectors form a basis of the vector space on which \boldsymbol{C} acts (eg, they are orthogonal).

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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.

III. SINGULAR VALUE DECOMPOSITION

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

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

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The number of singular values is equal to the rank of X.

The rank of a matrix measures its *non-degeneracy*.

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For a general SVD, the columns of U are the eigenvectors of XX^T , and the columns of V are the eigenvectors of X^TX .

Also, the singular values of X are the square roots of the eigenvalues of XX^T and X^TX .

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NOTE

If data is centered, these are covariance matrices.

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Here "best" refers to the representation that minimizes the squared orthogonal distances from the points to the subspace.

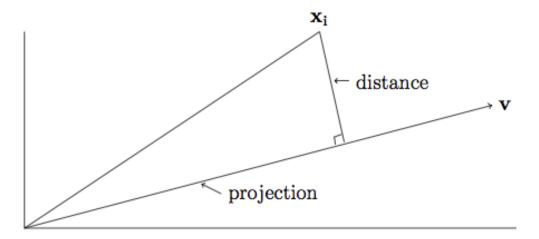


Figure 4.1: The projection of the point $\mathbf{x_i}$ onto the line through the origin in the direction of \mathbf{v}

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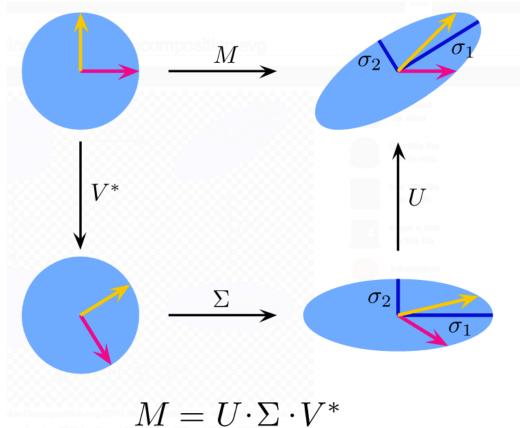
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The singular vectors of \mathcal{T} correspond to the lengths of the axes of the \mathcal{C} -dimensional ellipsoid.

The singular values give the magnitudes of the projection of each column of the original dataset on the elements of the new basis.

SINGULAR VALUE DECOMPOSITION



source: http://en.wikipedia.org/wiki/Singular_value_decomposition

SINGULAR VALUE DECOMPOSITION

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 More numerically stable and can be more efficient to calculate (than PCA)

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Latent semantic analysis, etc.

III. OTHER METHODS

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FACTOR ANALYSIS

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

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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:

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

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In practice, PCA is often used for factor analysis, after modifying the covariance matrix somewhat. But it can also allow for non-isotropic errors, and there are other methods for fitting as well, and different theoretical concerns.

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

But sometimes linear techniques are not sufficient.

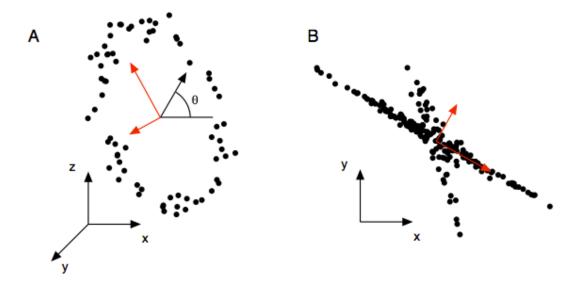
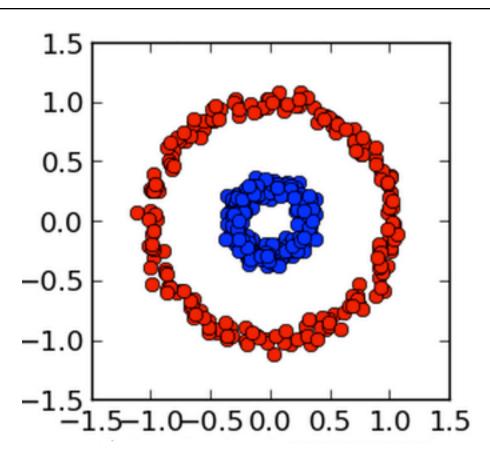


FIG. 6 Example of when PCA fails (red lines). (a) Tracking a person on a ferris wheel (black dots). All dynamics can be described by the phase of the wheel θ , a non-linear combination of the naive basis. (b) In this example data set, non-Gaussian distributed data and non-orthogonal axes causes PCA to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



multidimensional scaling: low-dim embedding that preserves pairwise distances

locally linear embedding: approximates local structure of data (neighborhood preserving embedding)

See sklearn manifold

Some methods for nonlinear dimensional reduction (or *manifold learning*) include:

multidimensional scaling: low-dim embedding preserves pairwise distances

locally linear embedding: approximates local structure of data (neighborhood preserving embedding)

kernel PCA: exploits PCA dependence on inner product (same logic as SVM)

isomap: nonlinear dimension reduction via MDS using geodesic (surface-bound) distances

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See sklearn.decomposition and sklearn manifold

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See

sklearn.decomposition and sklearn.manifold



And more!

In any case, key difficulties with dimensionality reduction are time/space complexity, randomness (eg different results for different runs), and selecting the number of dimensions in the lower-dim subspace.

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Furthermore, there's an obvious (bias/variance) tradeoff involved with the number of subspace dimensions and the size of approximation error.

INTRO TO DATA SCIENCE

IV. EXERCISE