





## Clustering Text

Natural Language Processing

Some slide content based on textbooks:

Speech and Language Processing: An Introduction to Natural Language Processing, Computational Linguistics, and Speech Recognition by Daniel Jurafsky and James H. Martin

ALICE was beginning to get very tired of sitting by her six the bank, and of having nothing to do: or wice she had peeped into the book her six the book her six the bank, and of having nothing to do: or without pictures or combined to be a six that is the use of a book her six the bank, and the bank had been also be a six could, for the hot day may be a six could, for the hot day may be a six could, for the hot day may be a six could, for the hot day may be a six could, for the hot day and six could, for the hot day and six could, for the hot day and six could, for the hot day white rabbit with pink eyes ran cit by her was nothing so very much of the six that he was nothing so very much of the six that it has not six that he was nothing so very much of the six

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#### Lecture content:

- Why cluster text?
- Similarities between documents
- Basic clustering algorithms
- Evaluating clustering
- Topic Modelling
- Visualisation via dimensionality reduction

Why cluster text?

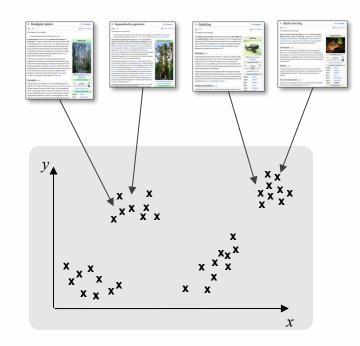
## What is text clustering?

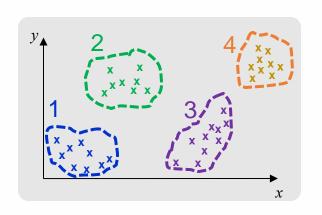
Process of grouping documents into coherent subgroups

• based on some distance function or similarity measure

For text documents, usually means grouping documents into:

- same **topic**: e.g. sports news vs politics vs technology vs ...
- same emotion: e.g. angry, disappointed, elated, hopeful, . . .
- same problem to be fixed: e.g. no heating (faulty thermostat), unable to print (missing drivers), no hot water, etc.
- same **complaint/praise**: e.g. hotel has beautiful view, motel on noisy road, breakfast was excellent, . . .
- same **author**: e.g. Shakespeare, Dylan, Eminem, Trump, Satoshi Nakamoto
- same **text generator**: e.g. used to generate spam emails
- same **source document**: e.g. for de-duplicating content during a web-crawl





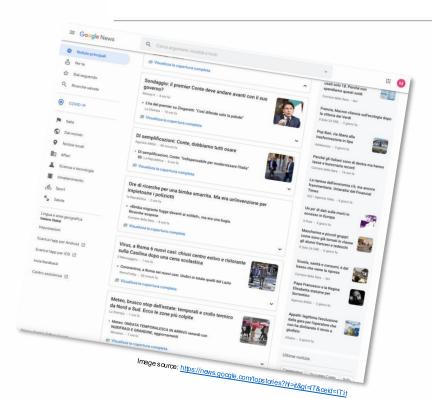
## Example Clustering Applications

News aggregators like Google News (<a href="https://news.google.com/">https://news.google.com/</a>) use clustering to summarise/characterise a collection:

- what **topics** are being discussed?
- how prevalent is each topic?
- what topics are trending?
- which articles are most exemplary (central to topic)?

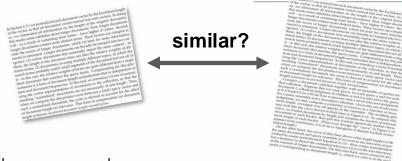
In security applications, such as phishing and malware detection,

- determine which groups might be malicious (based on a few identified instances)
- use clusters to discover useful **features** for describing documents
- what aspects distinguish the groups?



# Representing documents & measuring similarity

## Document similarity

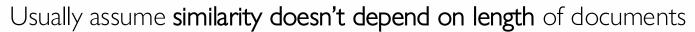


To cluster documents need to measure similarity between them

- based on the similarity of their content
- which is subjective of course: is Stephen King closer to Agatha Christie or J K Rowling?

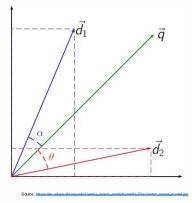
Traditionally represent documents as **tf-idf vectors** (scaled **bag-of-words**)

- sparse representation: most values in vector are zero
- so measure similarity based on **number of shared words**
- scaled by the importance of the word: how rare is it?



- so compute **cosine similarity** between vectors
- = dot product between length normalised vectors

$$sim(\mathbf{d}_1, \mathbf{d}_2) = \frac{\mathbf{d}_1 \cdot \mathbf{d}_2}{||\mathbf{d}_1|| ||\mathbf{d}_2||} \qquad ||\mathbf{x}|| = \sqrt{\frac{\mathbf{d}_1 \cdot \mathbf{d}_2}{||\mathbf{d}_1|| ||\mathbf{d}_2||}}$$



$$||\mathbf{x}|| = \sqrt{\sum_{i=1}^{n} x_i^2}$$

# Clustering techniques

## Types of clustering algorithms

Many different clustering algorithms in common use

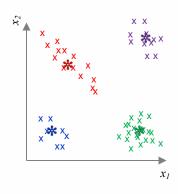
- k-Means/k-Medoids
- Agglomerative hierarchical clustering
- Density-based clustering
- Spectral clustering
- Topic Models

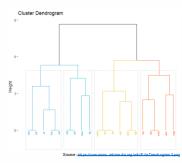
All these models are implemented in sklearn:

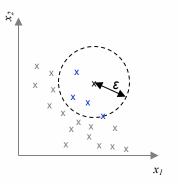
https://scikit-learn.org/stable/modules/clustering.html

We'll concentrate on those most used with text:

• primarily k-Means & Topic Models







## Clustering Algorithms: k-Means

## k-Means clustering

"Goto" clustering method

simple, fast and relatively robust

How does it work?

- searches for exactly *k* clusters
- represents each cluster by its centroid



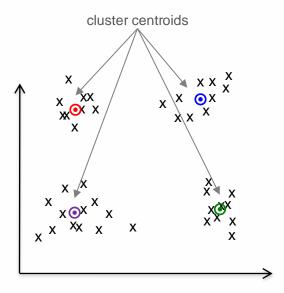
#### Advantages:

- scales well to large collections
- (requires no pairwise distance calculations)



#### **P** Disadvantages:

- searches for globular clusters
- implicitly assumes Euclidean distance metric (not ideal for text)
- number of clusters must be specified in advance



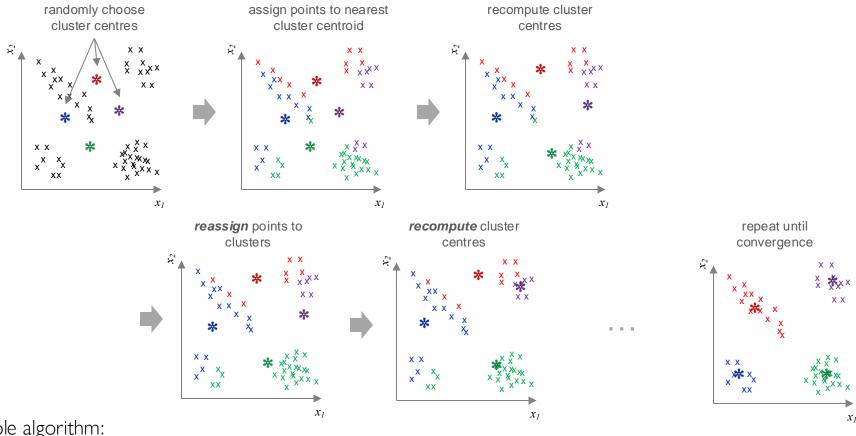
#### Centroid:

$$\vec{\mu}_C = \frac{1}{|C|} \sum_{\vec{x} \in C} \vec{x}$$

#### Euclidean distance:

$$d(x,y) = \sqrt{\sum_{i=1}^{m} (x_i - y_i)^2}$$

### k-Means clustering: the process



Simple algorithm:

- initialize k centroids randomly
- (re)assign each data point to closest centroid
- (re)compute centroids by averaging data points in cluster
- repeat steps (2) & (3) until convergence, when centroids stop moving

algorithm minimises variance of clusters (squared distances between points & cluster centres)

$$\min \sum_{i=1}^{k} \sum_{\vec{x} \in C_i} d(\vec{x}, \vec{\mu}_{C_i})^2$$

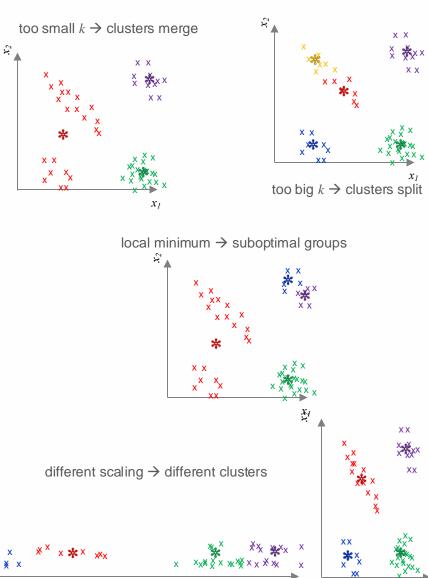
## k-Means clustering — potential problems

Potential problems with k-means:

- choosing "right" value of k is critical
  - can use elbow method to choose k
- algorithm can converge on local minimum
  - run algorithm multiple times algorithm

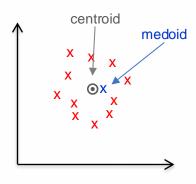
As for all clustering algorithms, scaling effects similarity measure

- and thus clusters found
- but for text have both tf-idf weighting and document length normalisaton



## Clustering Algorithms: k-medoids

#### k-medoids



Represents each cluster by its **medoid** rather than its centroid

- medoid is one of the datapoints from dataset
  - point that is closest (*smallest average distance*) to all other points
- at each iteration, algorithm must:
  - reassign datapoints to cluster with closest medoid, and then recompute medoids



#### Advantage:

- clustering algorithm can be used with other metrics, not just Euclidean
- medoid is a real document so provides realistic representation of cluster



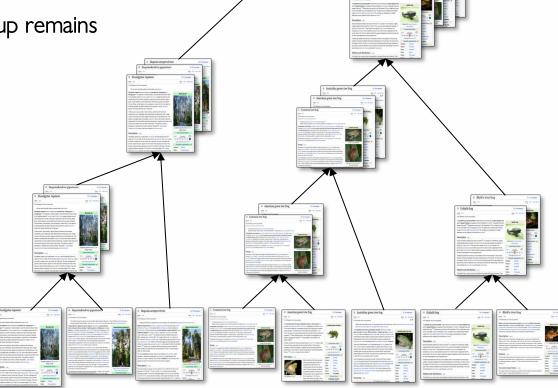
#### Disadvantage:

- algorithm has much higher computational complexity with respect to k-means
- needs to calculate distances for pairs of datapoints, which is an  $O(n^2)$  operation

# Clustering Algorithms: Agglomerative Hierarchical Clustering

Hierarchical Agglomerative Clustering Hierarchical clustering builds *hierarchy* of clusters called a **dendrogram** agglomerative clustering builds clusters by merging groups in bottom-up fashion

- assign each document to its own group
- merge 2 groups that are most similar
- repeat until only one group remains



## Agglomerative clustering: linking

Hierarchical clustering needs to compute distances between groups of documents

- uses a linkage criteria:
  - complete-linkage: **maximum** distance between points in 2 groups
  - single-linkage: **minimum** distance across groups
  - average-linkage: average distance across groups
- note: linking criteria used influences shape of clusters found
  - complete- or average-linkage will restrict to tight, globular clusters
  - single-linkage clustering will allow for finding long, thin clusters



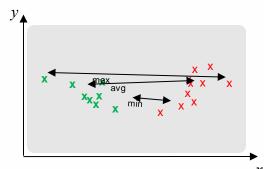
#### Advantage:

- works with any distance metric / similarity function
- dendrogram provides information about underlying structure of data



#### Disadvantage:

high time complexity makes it unsuitable for large datasets



# Clustering Algorithms: DBScan

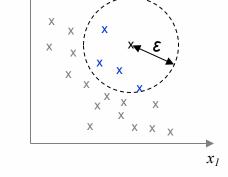
#### DBScan: what is it?

Density-Based Spatial Clustering of Applications with Noise

- #clusters not defined by user but inferred from data
- density-based algorithm: clusters found in high-density regions

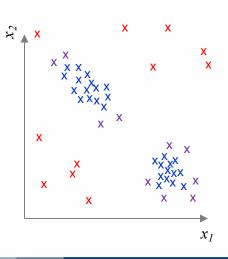
Algorithm takes 2 parameters:

- $\varepsilon$  = radius of **neighbourhood** around each point
- minPoints = minimum # of points required to form cluster

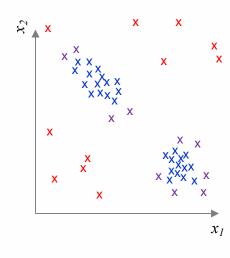


How does it work?

- # of datapoints in  $\varepsilon$ -neighbourhood is **density estimate**
- and minPoints is just a threshold on density
- algorithm classifies each point as either:
  - core point: has  $\geq$ minPoints neighbours inside radius  $\varepsilon$
  - border point: not core, but within  $\varepsilon$  of core point
  - noise point: neither core nor border point



## DBScan: pros and cons





#### Advantages:

- no need to choose number of clusters in advance
- identifies and ignores noise/outlier instances
- find arbitrarily (oddly) shaped clusters

#### Disadvantages:

- performance depends critically on chosen parameters (usually set  $\varepsilon$  as a function of average distance to *MinPoints* nearest neighbour)
- may not work well if clusters have different densities

## Evaluating Clustering

## Evaluating clustering









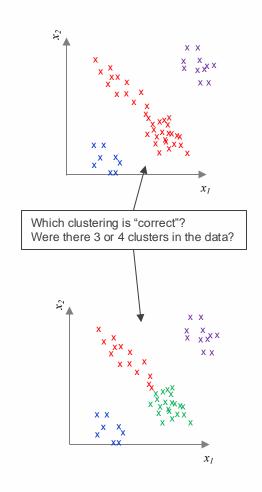




- usually don't have access to ground-truth labelling
- without it, evaluation can be subjective (people disagree on what "right" clusters are)

Two ways to evaluate results of clustering algorithm:

- intrinsic evaluation:
  - check how coherent and well separated clusters are
  - typical evaluation measures:
    - Within-cluster Sum of Squares (cluster variance)
    - Silhouette Coefficient: compares average distance to point in current cluster with points in next closest cluster
- 2. extrinsic evaluation:
  - check how well clusters align with known document labels
  - typical evaluation measures:
    - Homogeneity, Completeness and V-Measure: effectively Precision, Recall and F-measure applied to clustering problems

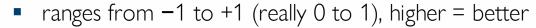


#### Evaluation: silhouette

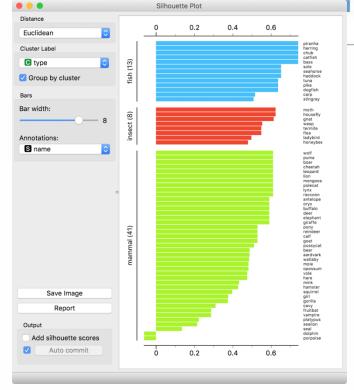
#### Silhouette coefficient

- score calculated **separately** for each sample
- uses distance metric (e.g., Euclidean)
- find average distance from sample  $x_i$  to all samples:
  - in same cluster (denote  $a_i$ )
  - in next closest cluster (denote  $b_i$ )
- compute coefficient as:

$$s_i = \frac{b_i - a_i}{\max(a_i, b_i)}$$



- average silhouette score over all datapoints to get single value representing quality of clustering
- often use silhouette plots to visualise quality of clusters
- issue: expensive to calculate, since done independently for each datapoint



Silhouette-plot-orange:

https://commons.wikimedia.org/wiki/File:Silhouette-plot-orange.png

## Evaluation: sum of squares

Very common to compute sum of squares values:

- Within-cluster Sum of Squares
  - sum over clusters the squared distances between cluster centroid and each point
- Between-cluster Sum of Squares
  - sum over clusters the squared distances between data centroid and each cluster centroid (weighted by data)
- Variance Ratio Criterion (aka Calinski-Harabasz index)
  - combines both measures into single metric
  - measures whether clusters are dense and well separated
  - faster to compute than Silhouette index, but really just objective function of k-Means

$$WSS(C) = \sum_{i=1}^{k} \sum_{x_j \in C_i} d(x_j, \mu_i)^2$$

BSS(C) = 
$$\sum_{i=1}^{k} |C_i| d(\mu, \mu_i)^2$$

$$s = \frac{BSS(C)}{WSS(C)} \times \frac{N-k}{k-1}$$

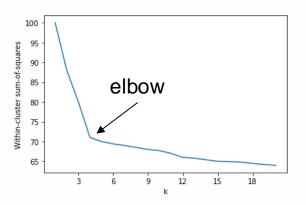
## Evaluating clustering — cont.

**Intrinsic** evaluation measures can be useful to:

- visualise cluster quality in Silhouette plots
- select a value for k using the Elbow method
  - plot within cluster sum of squares versus number of clusters
  - as k gets larger, intrinsic measure always improves
  - so don't look for best value, but rather biggest jump:
    - point where performance increases substantially, but then slows is called **elbow**



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# Topic Modelling

## What are topic models?

#### **Soft clustering** of documents

allows documents to belong to multiple clusters

#### Provides low-dimensional representation of documents

compared to high dimensional vocabulary space

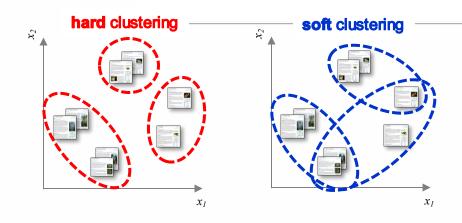
Each cluster referred to as a "topic"

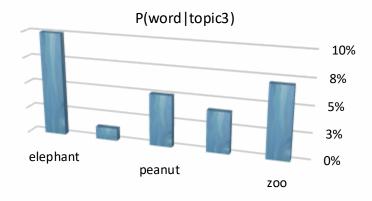
- described by distribution over words
- terms should be coherent (discuss single theme)

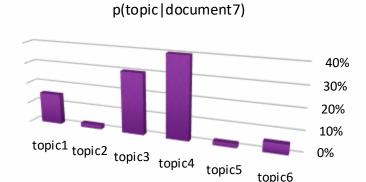
Topic vector = probability distribution over words

$$P(\texttt{elephant}|\texttt{topic3}) = 0.09$$

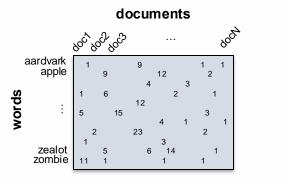
Document vector = probability distribution over topics P(topic3|doc7) = 0.1

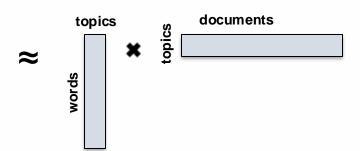






## Matrix Decomposition





Topic Modeling is a form of matrix decomposition

- decomposes terms\*documents count matrix
- into 2 smaller matrices: terms\*topics and topics\*documents

Much smaller number of parameters need to be estimated: V\*T+T\*D <<< V\*D

• for vocabulary size V, document count D and topic count T

Most famous technique is Latent Dirichlet Allocation (LDA)

- so named because it uses a Dirichlet prior when estimating the parameters
- related to Non-negative Matrix Factorization (NMF)
- also Latent Semantic Indexing (LSI), which applies Singular Value Decomposition (SVD) to TFIDF matrix

## What is topic modelling useful for?

#### Factorization/clustering generalizes observed term counts:

- making document representations more dense & useful
- allows calculating more meaningful distances between documents

#### Deals with problems of:

- polysemy: bank (financial institution or edge of river?)
- synonymy: cancer = oncology ?
- short documents: limited vocabulary

#### Sometimes useful for visualizing collections

• after further dimensionality reduction (using say t-SNE)



Image source: https://www.flickr.com/photos/wwarby/4915969081

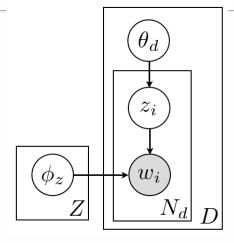
## Details of LDA generative model

#### Generative model for latent Dirichlet Allocation (LDA)

- 1. choose word proportions for each topic
- 2. choose topic proportions for each document
- 3. for each position in document
  - 1. choose **topic** based on document proportions
  - 2. choose word based on topic proportions

#### Estimating parameters for topic model:

- requires iteratively updating parameters
  - 1. topic probabilities for each document
  - 2. word probabilities for each topic
- place Bayesian priors on parameters to avoid over-fitting
- use (Gibbs) sampling techniques to avoid local maxima
- hyperparameters  $\mathbf{a}$  and  $\boldsymbol{\beta}$  determine the prior on the topic/document distributions (how concentrated they are)

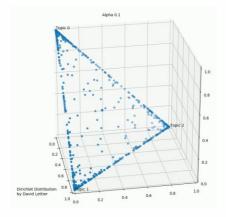


 $\phi_z \sim Dirichlet(\beta)$ 

 $\theta_d \sim Dirichlet(\alpha)$ 

 $z_i \sim Discrete(\theta_d)$ 

 $w_i \sim Discrete(\phi_{z_i})$ 

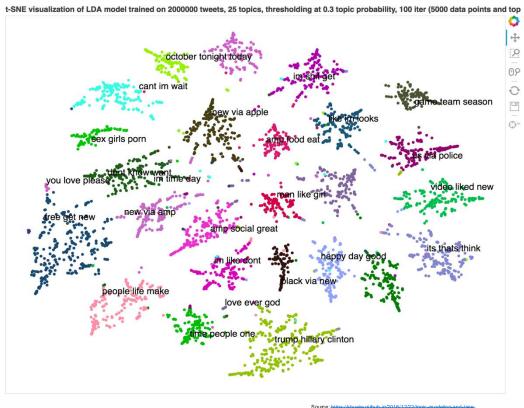


Source: David Lettier https://medium.com/@lettier/how-does-lda-work-illexplain-using-emoji-108abf40fa7d

## Applications of Topic Models

Useful for visualising collections

- represent topics by most frequent terms
- common also to show changes to collection over time



Source: https://shuaiw.gi.thub.io/2016/12/22/topic-modeling-and-duisus/artion.html

# Visualising document vectors: dimensionality reduction techniques

# Can project high-dimensional data into lower dimensions using:

### 1. a linear projection

with Principle Component Analysis

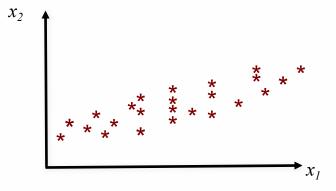
#### 2. a non-linear projection

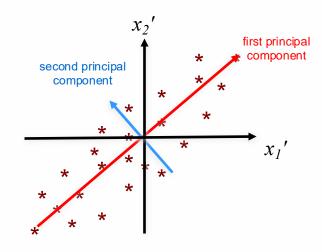
e.g. using t-distributed Stochastic Neighbor Embeddings

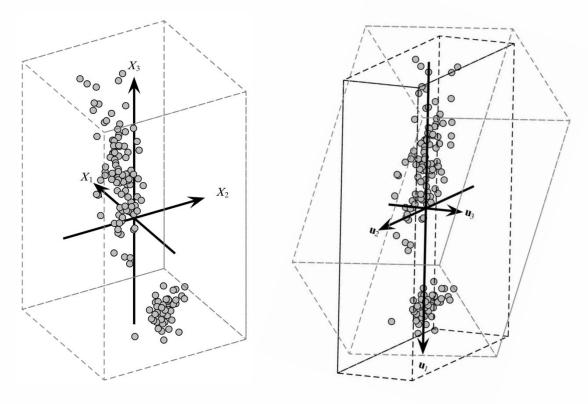
## Principal Component Analysis (PCA)

Investigates main sources of variation in dataset

- by finding **projection** that captures **largest** amount of variation in data
- method:
  - 1. scale features to have zero mean, unit variance
  - 2. find **direction** in **d**-dimensions that represent largest variation in data (k < d)
  - 3. remove variation from data
  - 4. repeat until k dimensions found
- use new dimensions to approximate original data using only largest components







#### For more information on PCA

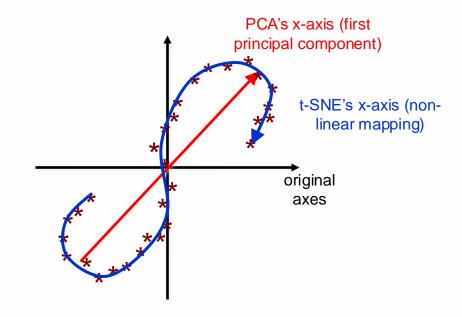
- see Chapter 7 of the textbook "Data Mining and Analysis"
- and: <a href="http://sebastianraschka.com/Articles/2015">http://sebastianraschka.com/Articles/2015</a> pca in 3 steps.html

# t-distributed Stochastic Neighbor Embedding

### Non-linear Dimensionality Reduction

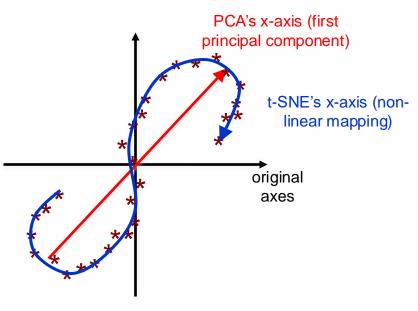
Data in high dimensions never fills the entire space

 always lives on some lowerdimensional manifold



t-SNE is a non-linear dimensionality reduction technique

- used to map high-dimensional data into 2 or 3 dimensions
  - points from original space mapped onto "map points" in 2D/3D
- unlike PCA:
  - mapped point is **not linear combination** of **original attribute values**
  - axis of mapped space is not linear combination (rotation) of original axes



#### t-SNE:

- tries to preserve local distances to nearby points
  - unlike PCA which tries to preserve global (long range) distances between points
- converts distances between data points to joint probabilities
  - then maps original points to lower dimensional points such that position of mapped points preserves structure of the data
    - i.e. similar data points are assigned nearby map points
    - while dissimilar data points are assigned distant map points

## t-SNE: algorithm

#### The t-SNE algorithm has two main steps:

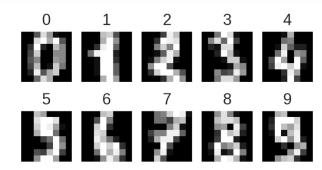
- 1. define probability distribution over pairs of high-dimensional points:
  - such that pairs of similar data points have high probability of being picked
  - and dissimilar points have extremely low probability of being picked
- 2. define a similar distribution over the points in the map space
  - and move around the mapped points to minimize Kullback—Leibler divergence between the two distributions
  - gradient descent is used to minimize the score

#### Caveats of t-SNE:

- perplexity parameter controls balance between local and global aspects of the data
- different initializations will lead to different results
- applies only to data with "reasonable" number of dimensions (e.g. 30-50)
  - if data has more dimensions, computational time will be prohibitive

### t-SNE example: handwritten digit dataset

- Handwritten Digits Data Set from the UCI machine learning repository <u>http://archive.ics.uci.edu/ml/datasets/Optical+Recognition+of+Handwritten+Digits</u>
- Contains 1797 images with 8x8 pixels each



.9

https://github.com/oreillymedia/t-SNE-tutorial

## Conclusions

#### Conclusions

- Many applications of text clustering
- Many clustering techniques can be applied to tf-idf vectors
- Evaluating clustering can be subjective
- Topic modeling builds lower-dimensional representation of documents by performing soft clustering
- Dimensionality reduction techniques can be used to visualize document vectors