Clustering, K-Means, and K-Nearest Neighbors

CMSC 678 UMBC

Recap from last time...

Geometric Rationale of LDiscA & PCA

Objective: to rigidly rotate the axes of the D-dimensional space to new positions (principal axes):

ordered such that principal axis 1 has the highest variance, axis 2 has the next highest variance,, and axis D has the lowest variance

covariance among each pair of the principal axes is zero (the principal axes are uncorrelated)

L-Dimensional PCA

1. Compute mean μ , priors, and common covariance Σ

$$\Sigma = \frac{1}{N} \sum_{i: v_i = k} (x_i - \mu)(x_i - \mu)^T \qquad \qquad \mu = \frac{1}{N} \sum_i x_i$$

- 2. Sphere the data (zero-mean, unit covariance)
- Compute the (top L) eigenvectors, from sphere-d data, via V

$$X^* = VD_BV^T$$

4. Project the data

Outline

Clustering basics

K-means: basic algorithm & extensions

Cluster evaluation

Non-parametric mode finding: density estimation

Graph & spectral clustering

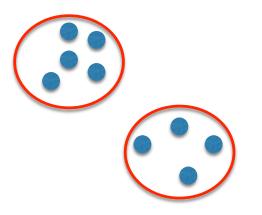
Hierarchical clustering

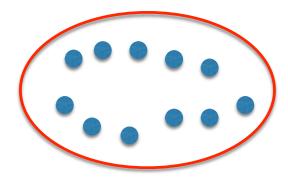
K-Nearest Neighbor

Clustering

Basic idea: group together similar instances

Example: 2D points

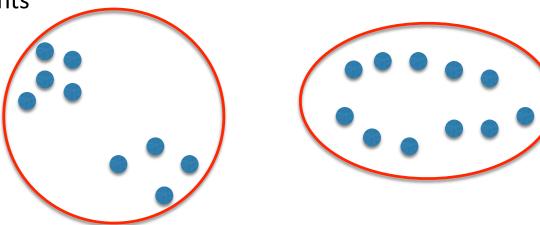




Clustering

Basic idea: group together similar instances

Example: 2D points



One option: small Euclidean distance (squared)

$$\operatorname{dist}(\mathbf{x}, \mathbf{y}) = ||\mathbf{x} - \mathbf{y}||_2^2$$

Clustering results are crucially dependent on the measure of similarity (or distance) between points to be clustered

Clustering algorithms

Simple clustering: organize elements into k groups

K-means

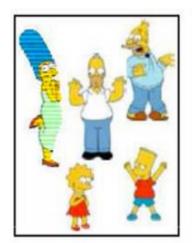
Mean shift

Spectral clustering

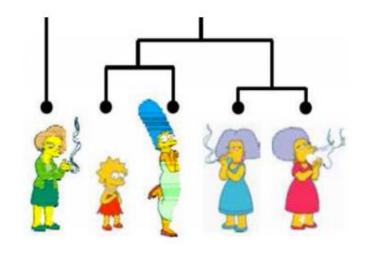
Hierarchical clustering: organize elements into a hierarchy

Bottom up - agglomerative

Top down - divisive



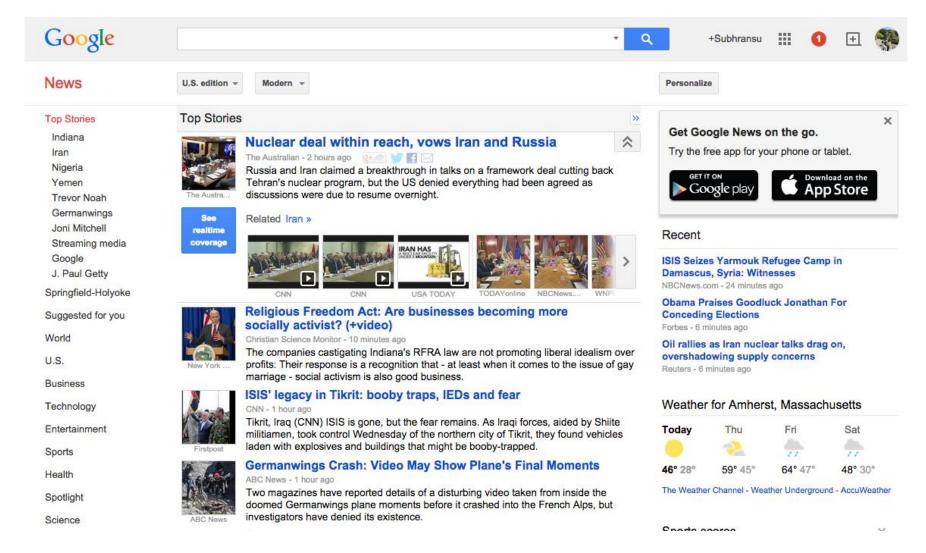




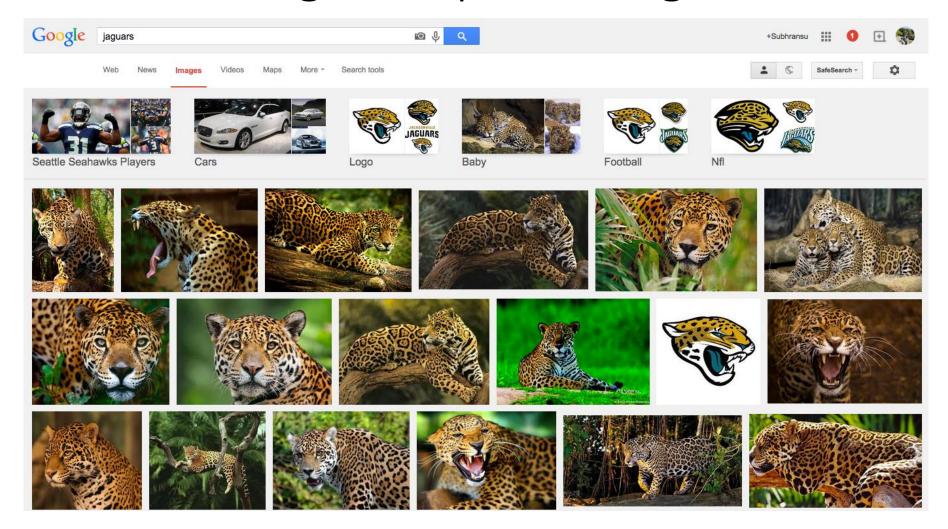
Clustering examples: Image Segmentation



Clustering examples: News Feed



Clustering examples: Image Search



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Clustering using k-means

Data: D-dimensional observations $(x_1, x_2, ..., x_n)$

Goal: partition the **n** observations into **k** (\leq **n**) sets **S** = {S₁, S₂, ..., S_k} so as to minimize the within-cluster sum of squared distances

$$\arg\min_{\mathbf{S}} \sum_{i=1}^{k} \sum_{\mathbf{x} \in S_i} ||\mathbf{x} - \mu_i||^2$$
 cluster center

Lloyd's algorithm for k-means

Initialize k centers by picking k points randomly among all the points

Repeat till convergence (or max iterations)

Assign each point to the nearest center (assignment step)

$$\arg\min_{\mathbf{S}} \sum_{i=1}^{k} \sum_{\mathbf{x} \in S_i} ||\mathbf{x} - \mu_i||^2$$

Estimate the mean of each group (update step)

Properties of the Lloyd's algorithm

Guaranteed to converge in a finite number of iterations objective decreases monotonically local minima if the partitions don't change.

finitely many partitions → k-means algorithm must converge

Running time per iteration

Assignment step: O(NKD)

Computing cluster mean: O(ND)

Issues with the algorithm:

Worst case running time is super-polynomial in input size No guarantees about global optimality

Optimal clustering even for 2 clusters is NP-hard [Aloise et al., 09]

k-means++ algorithm

A way to pick the good initial centers

Intuition: spread out the k initial cluster centers

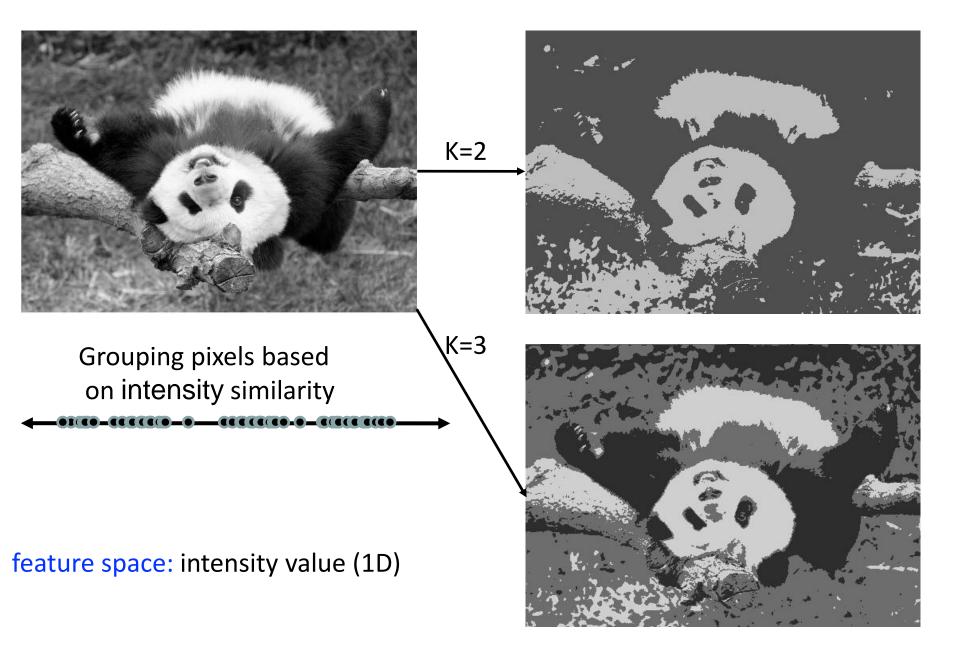
The algorithm proceeds normally once the centers are initialized

[Arthur and Vassilvitskii'07] The approximation quality is O(log k) in expectation

k-means++ algorithm for initialization:

- 1. Chose one center uniformly at random among all the points
- 2. For each point **x**, compute D(**x**), the distance between x and the nearest center that has already been chosen
- 3. Chose one new data point at random as a new center, using a weighted probability distribution where a point **x** is chosen with a probability proportional to D(**x**)²
- 4. Repeat Steps 2 and 3 until k centers have been chosen

k-means for image segmentation



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Clustering Evaluation

(Classification: accuracy, recall, precision, F-score)

Greedy mapping: one-to-one

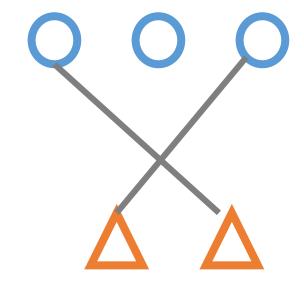
Optimistic mapping: many-to-one

Rigorous/information theoretic: V-measure

Clustering Evaluation: One-to-One

Each modeled cluster can at most only map to one gold tag type, and vice versa

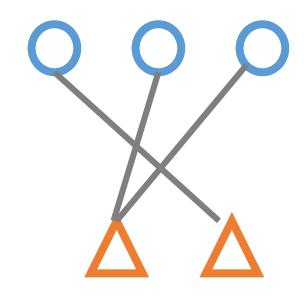
Greedily select the mapping to maximize accuracy



Clustering Evaluation: Many (classes)-to-One (cluster)

Each modeled cluster can map to at most one gold tag types, but multiple clusters can map to the same gold tag

For each cluster: select the majority tag



Rosenberg and Hirschberg (2008): harmonic mean of homogeneity and completeness

$$H(X) = -\sum_{i} p(x_i) \log p(x_i)$$
entropy

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entropy

Rosenberg and Hirschberg (2008): harmonic mean of *homogeneity* and *completeness*

k → cluster c → gold class

Homogeneity: how well does each gold class map to a single cluster?

"In order to satisfy our homogeneity criteria, a clustering must assign only those datapoints that are members of a single class to a single cluster. That is, the class distribution within each cluster should be skewed to a single class, that is, zero entropy."

homogeneity =
$$\begin{cases} 1, & H(K,C) = 0 \\ 1 - \frac{H(C|K)}{H(C)}, & \text{o/w} \end{cases}$$

relative entropy is maximized when a cluster provides no new info. on class grouping \rightarrow not very homogeneous

Rosenberg and Hirschberg (2008): harmonic mean of *homogeneity* and *completeness*

k → cluster c → gold class

Completeness: how well does each learned cluster cover a single gold class?

"In order to satisfy the completeness criteria, a clustering must assign all of those datapoints that are members of a single class to a single cluster. "

completeness =
$$\begin{cases} 1, & H(K,C) = 0 \\ 1 - \frac{H(K|C)}{H(K)}, & \text{o/w} \end{cases}$$

relative entropy is maximized when each class is represented uniformly (relatively) → not very complete

Rosenberg and Hirschberg (2008): harmonic mean of *homogeneity* and *completeness*

k → cluster c → gold class

Homogeneity: how well does each gold class map to a single cluster?

homogeneity =
$$\begin{cases} 1, & H(K,C) = 0 \\ 1 - \frac{H(C|K)}{H(C)}, & \text{o/w} \end{cases}$$

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Rosenberg and Hirschberg (2008): harmonic mean of *homogeneity* and *completeness*

 a_{ck} = # elements of class c in cluster k

Homogeneity: how well does each gold class map to a single cluster?

Completeness: how well does each learned cluster cover a *single* gold class?

homogeneity =
$$\begin{cases} 1, & H(K,C) = 0 \\ 1 - \frac{H(C|K)}{H(C)}, & \text{o/w} \end{cases}$$

$$H(C|K) = -\sum_{k}^{K} \sum_{c}^{C} \frac{a_{ck}}{N} \log \frac{a_{ck}}{\sum_{c'} a_{c'k}}$$

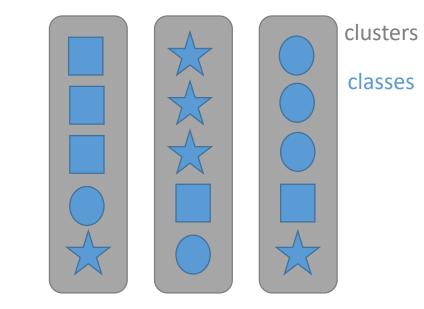
$$H(K|C) = -\sum_{c}^{C} \sum_{k}^{K} \frac{a_{ck}}{N} \log \frac{a_{ck}}{\sum_{k'} a_{ck'}}$$

completeness =
$$\begin{cases} 1, & H(K,C) = 0 \\ 1 - \frac{H(K|C)}{H(K)}, & \text{o/w} \end{cases}$$

Rosenberg and Hirschberg (2008): harmonic mean of homogeneity and completeness

Homogeneity: how well does each gold class map to a single cluster?

Completeness: how well does each learned cluster cover a *single* gold class?



	K C	
H(C K) =	$-\sum\sum \frac{a_{ck}}{N}$ lo	a_{ck}
H(C K) = -	$\underset{k}{\overset{\sim}{\smile}} \underset{c}{\overset{\sim}{\smile}} N$	$\sum_{c'} a_{c'k}$

$$H(K|C) = -\sum_{C}^{C} \sum_{k}^{K} \frac{a_{ck}}{N} \log \frac{a_{ck}}{\sum_{k} a_{ck}}$$

a _{ck}	K=1	K=2	K=3
	3	1	1
	1	1	3
\Rightarrow	1	3	1

Homogeneity = Completeness = V-Measure=0.14

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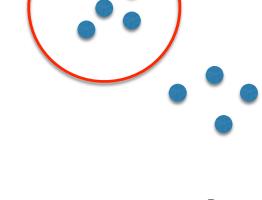
K-Nearest Neighbor

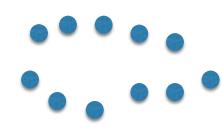
Clustering using density estimation

One issue with k-means is that it is sometimes hard to pick k

The mean shift algorithm seeks modes or local maxima of density in the feature space

Mean shift automatically determines the number of clusters





$$K(\mathbf{x}) = \frac{1}{Z} \sum_{i} \exp\left(-\frac{||\mathbf{x} - \mathbf{x}_{i}||^{2}}{h}\right)$$

Kernel density estimator

Small h implies more modes (bumpy distribution)

Mean shift algorithm

For each point x_i :

find m_i, the amount to shift each point x_i to its centroid

return {m_i}

Mean shift algorithm

```
For each point x_i:

set m_i = x_i

while not converged:

compute weighted average of neighboring point

return \{m_i\}
```

Mean shift algorithm

For each point
$$x_i$$
: Neighbors of x_i

set $m_i = x_i$

while not converged:

compute $m_i = \frac{\sum_{x_j \in N(x_i)} x_j K(m_i, x_j)}{\sum_{x_j \in N(x_i)} K(m_i, x_j)}$

return $\{m_i\}$

weighted average

self-clustering to based on kernel (similarity to other points)

Pros:

Does not assume shape on clusters

Generic technique

Finds multiple modes

Parallelizable

Cons:

Slow: O(DN²) per iteration
Does not work well for
high-dimensional
features

Mean shift clustering results









http://www.caip.rutgers.edu/~comanici/MSPAMI/msPamiResults.html

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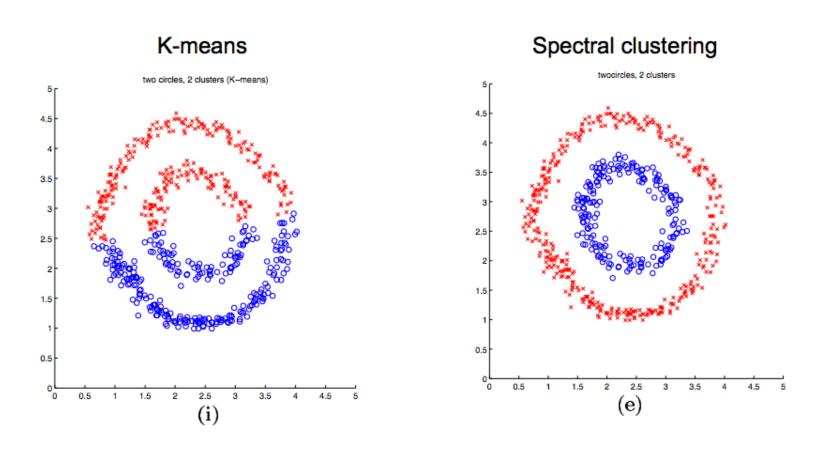
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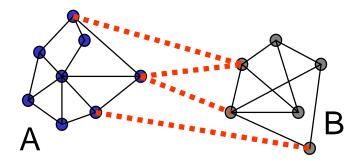
Spectral clustering



[Shi & Malik '00; Ng, Jordan, Weiss NIPS '01]

Spectral clustering

Group points based on the links in a graph



How do we create the graph?

Weights on the edges based on similarity between the points A common choice is the Gaussian kernel

$$W(i,j) = \exp\left(-\frac{||\mathbf{x}_i - \mathbf{x}_j||^2}{2\sigma^2}\right)$$

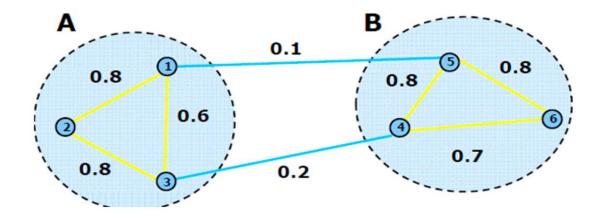
One could create

A fully connected graph

k-nearest graph (each node is connected only to its k-nearest neighbors)

Graph cut

Consider a partition of the graph into two parts A and B



Cut(A, B) is the weight of all edges that connect the two groups

$$\operatorname{Cut}(A,B) = \sum_{i \in A, j \in B} W(i,j) = 0.3$$

An intuitive goal is to find a partition that minimizes the cut min-cuts in graphs can be computed in polynomial time

Problem with min-cut

The weight of a cut is proportional to number of edges in the cut; tends to produce small, isolated components.

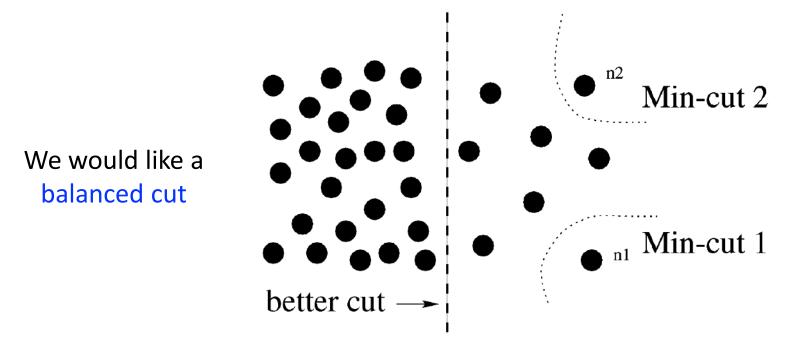


Fig. 1. A case where minimum cut gives a bad partition. [Shi & Malik, 2000 PAMI]

Graphs as matrices

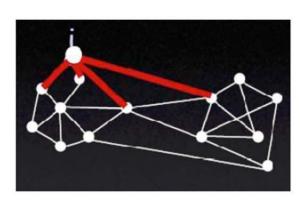
Let *W(i, j)* denote the matrix of the edge weights

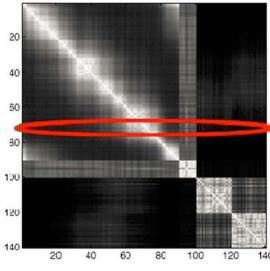
The degree of node in the graph is:

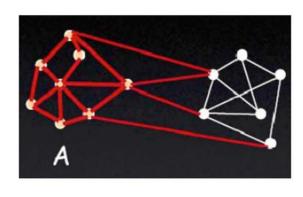
$$d(i) = \sum_{j} W(i, j)$$

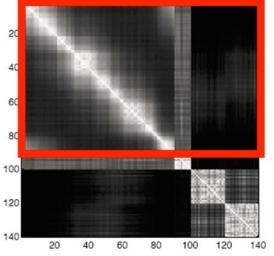
The volume of a set A is defined as:

$$Vol(A) = \sum_{i \in A} d(i)$$









Normalized cut

the connectivity between the groups relative to the volume of each group:

$$NCut(A, B) = \frac{Cut(A, B)}{Vol(A)} + \frac{Cut(A, B)}{Vol(B)}$$

$$\operatorname{NCut}(A, B) = \operatorname{Cut}(A, B) \left(\frac{\operatorname{Vol}(A) + \operatorname{Vol}(B)}{\operatorname{Vol}(A)\operatorname{Vol}(B)} \right)$$

minimized when Vol(A) = Vol(B)

a balanced cut

Minimizing normalized cut is NP-Hard even for planar graphs [Shi & Malik, 00]

Solving normalized cuts

W: the similarity matrix

D: a diagonal matrix with D(i,i) = d(i) — the degree of node i

y: a vector
$$\{1, -b\}^N$$
, $y(i) = 1 \leftrightarrow i \in A$ allow for differing penalty

The matrix (D-W) is called the Laplacian of the graph

$$\min_{\mathbf{x}} \text{NCut}(\mathbf{x}) = \min_{\mathbf{y}} \frac{\mathbf{y}^T (D - W)\mathbf{y}}{\mathbf{y}^T D\mathbf{y}}$$

subject to:
$$\mathbf{y}^T D \mathbf{1} = 0$$

 $\mathbf{y}(i) \in \{1, -b\}$

Solving normalized cuts

Normalized cuts objective:
$$\min_{\mathbf{x}} \mathrm{NCut}(\mathbf{x}) = \min_{\mathbf{y}} \frac{\mathbf{y}^T (D - W) \mathbf{y}}{\mathbf{y}^T D \mathbf{y}}$$

subject to:
$$\mathbf{y}^T D \mathbf{1} = 0$$

 $\mathbf{y}(i) \in \{1, -b\}$

Relax the integer constraint on **y**:

$$\min_{\mathbf{y}} \mathbf{y}^T (D - W) \mathbf{y}$$
; subject to: $\mathbf{y}^T D \mathbf{y} = 1, \mathbf{y}^T D \mathbf{1} = 0$

Same as: $(D-W)\mathbf{1} = 0$ (Generalized eigenvalue problem)

 $(D-W)\mathbf{y}=\lambda D\mathbf{y}$ \rightarrow the first eigenvector is \mathbf{y}_1 = 1, with the corresponding eigenvalue of 0

The eigenvector corresponding to the second smallest eigenvalue is the solution to the relaxed problem



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Hierarchical clustering

Agglomerative: a "bottom up" approach where elements start as individual clusters and clusters are merged as one moves up the hierarchy

Divisive: a "top down" approach where elements start as a single cluster and clusters are split as one moves down the hierarchy

Agglomerative clustering

Agglomerative clustering:

First merge very similar instances
Incrementally build larger clusters out
of smaller clusters

Algorithm:

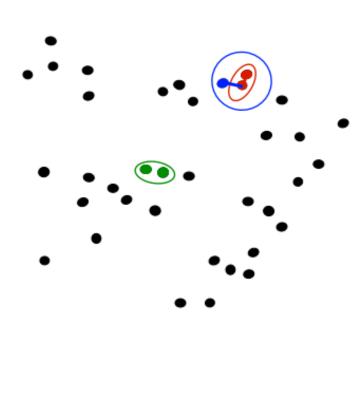
Maintain a set of clusters
Initially, each instance in its own cluster
Repeat:

Pick the two "closest" clusters

Merge them into a new cluster

Stop when there's only one cluster left

Produces not one clustering, but a family of clusterings represented by a dendrogram





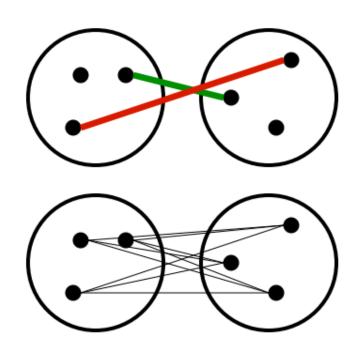
Agglomerative clustering

How should we define "closest" for clusters with multiple elements?

Closest pair: single-link clustering

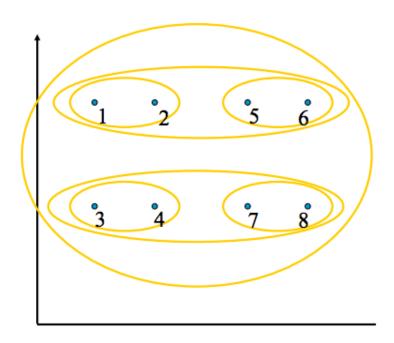
Farthest pair: complete-link clustering

Average of all pairs

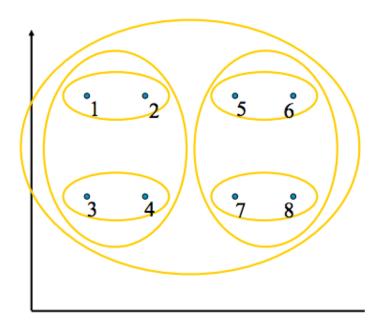


Agglomerative clustering

Closest pair (single-link clustering)



Farthest pair (complete-link clustering)



[Pictures from Thorsten Joachims]

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Nearest neighbor classifier

Will Alice like the movie?

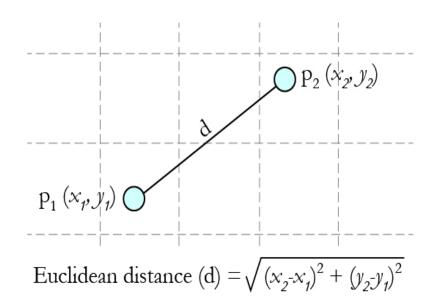
Alice and James are similar

James likes the movie \rightarrow

Alice must/might also like the movie

Represent data as vectors of feature values

Find closest (Euclidean norm) points



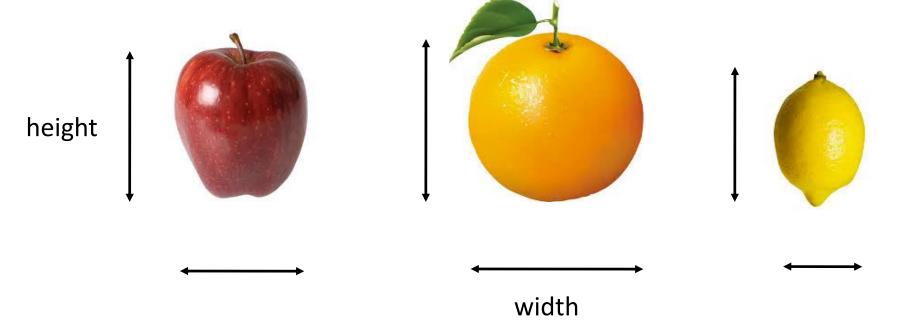
Nearest neighbor classifier

Training data is in the form of $(\mathbf{x}_1,y_1),(\mathbf{x}_2,y_2),\ldots,(\mathbf{x}_n,y_n)$

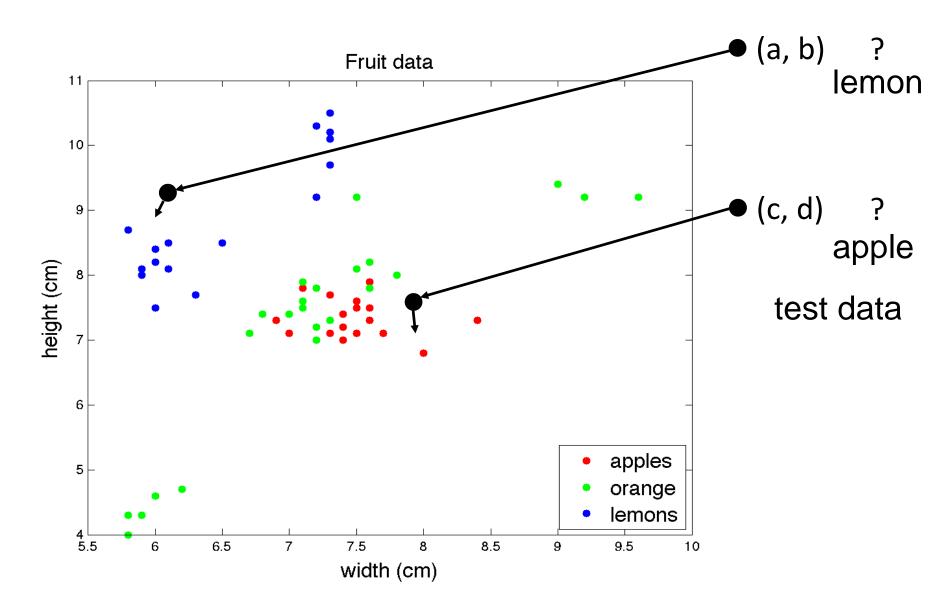
Fruit data:

label: {apples, oranges, lemons}

attributes: {width, height}

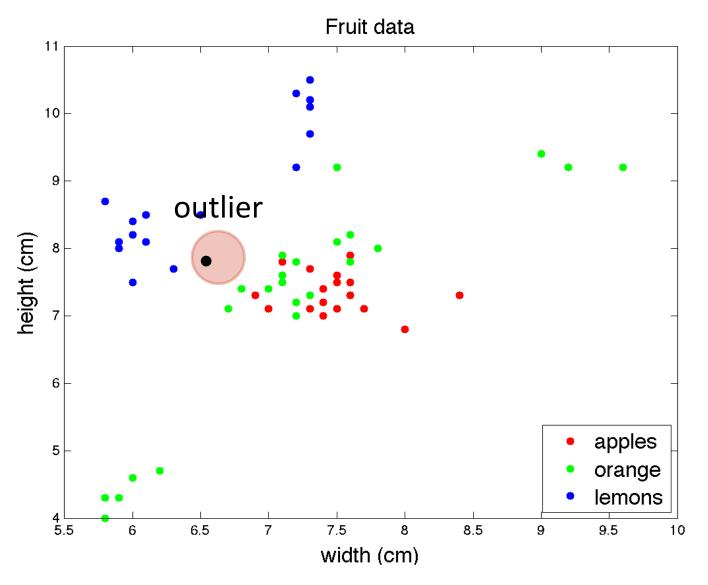


Nearest neighbor classifier



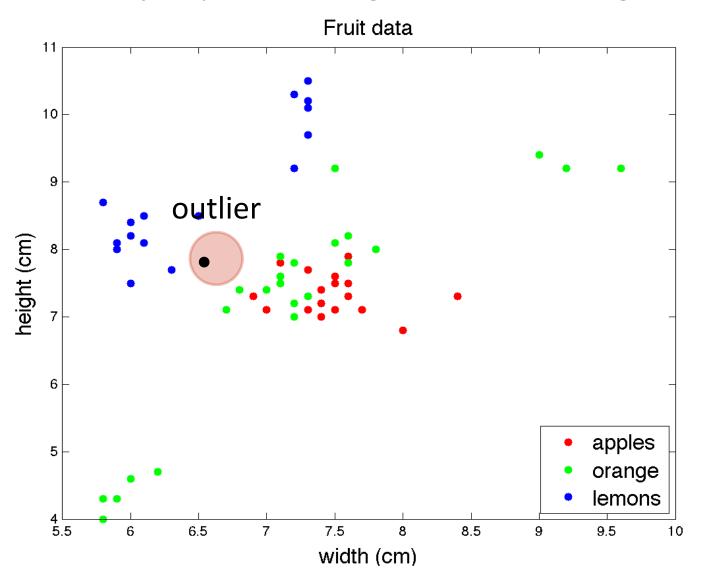
k-Nearest neighbor classifier

Take majority vote among the k nearest neighbors



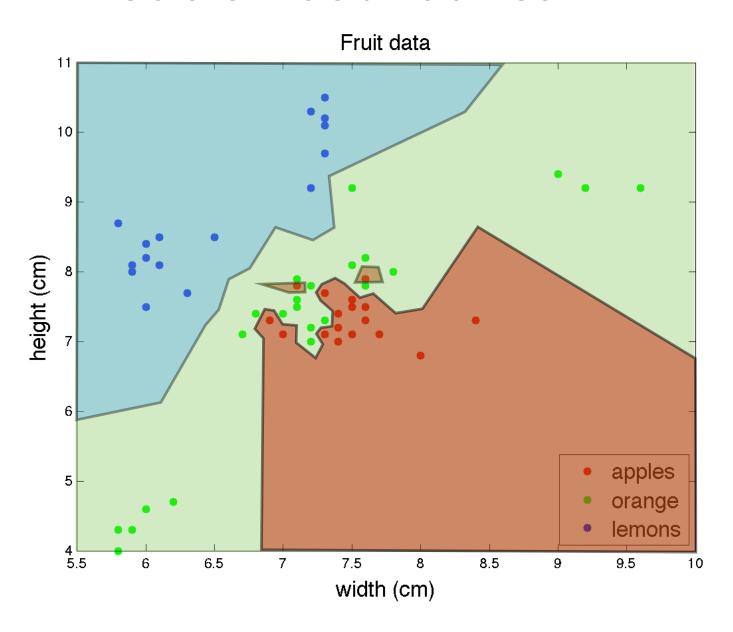
k-Nearest neighbor classifier

Take majority vote among the k nearest neighbors



What is the effect of k?

Decision boundaries: 1NN



Inductive bias of the kNN classifier

Choice of features

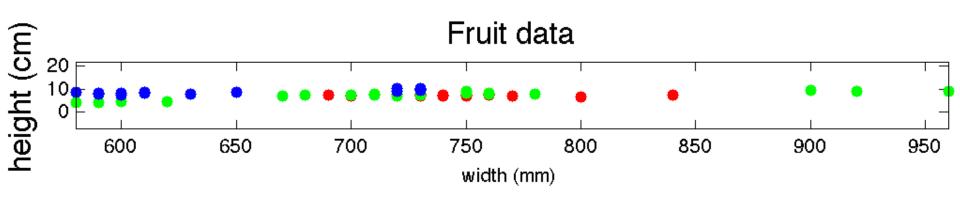
We are assuming that all features are equally important What happens if we scale one of the features by a factor of 100?

Choice of distance function

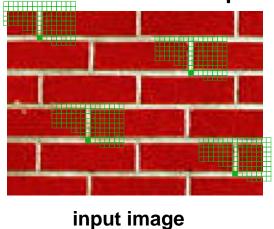
Euclidean, cosine similarity (angle), Gaussian, etc ...

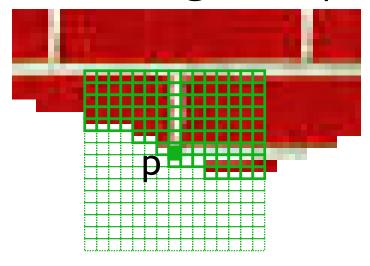
Should the coordinates be independent?

Choice of k



An example: Synthesizing one pixel





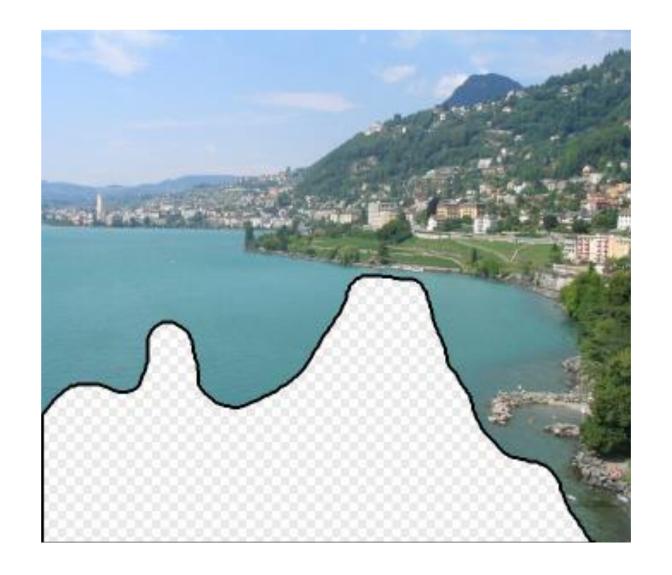
What is $P(\mathbf{x}|\text{neighborhood of pixels around }\mathbf{x})$

Find all the windows in the image that match the neighborhood

To synthesize **x**

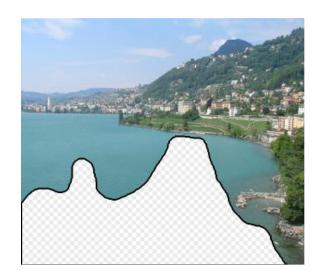
pick one matching window at random assign **x** to be the center pixel of that window

An **exact** match might not be present, so find the **best** matches using **Euclidean distance** and randomly choose between them, preferring better matches with higher probability



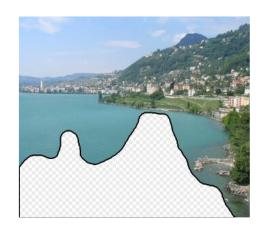
"Scene completion using millions of photographs", Hayes and Efros, TOG 2007

Nearest neighbors





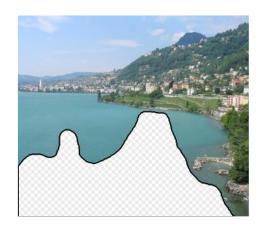
"Scene completion using millions of photographs", Hayes and Efros, TOG 2007







"Scene completion using millions of photographs", Hayes and Efros, TOG 2007







"Scene completion using millions of photographs", Hayes and Efros, TOG 2007

Practical issue when using kNN: speed

Time taken by kNN for N points of D dimensions

time to compute distances: O(ND)

time to find the k nearest neighbor

O(k N): repeated minima

O(N log N) : sorting

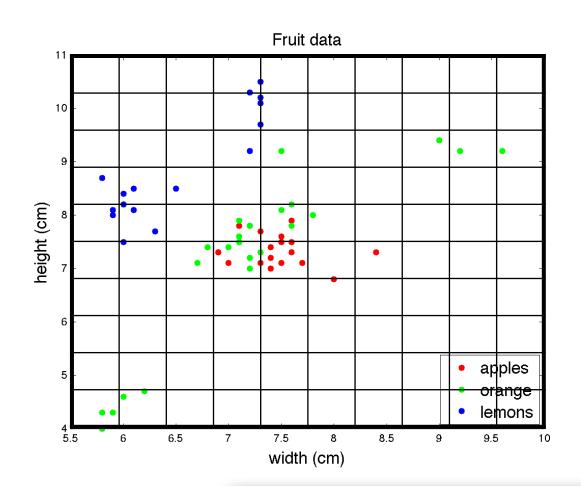
O(N + k log N): min heap

O(N + k log k): fast median

Total time is dominated by distance computation

We can be faster if we are willing to sacrifice exactness

Practical issue when using kNN: Curse of dimensionality



#bins =
$$10x10$$

d = 2

#bins =
$$10^{d}$$
 d = 1000

Atoms in the universe: $^{\sim}10^{80}$

How many neighborhoods are there?

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