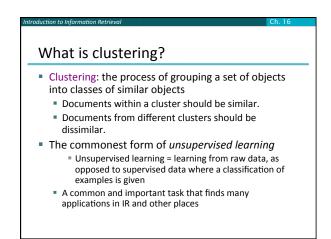
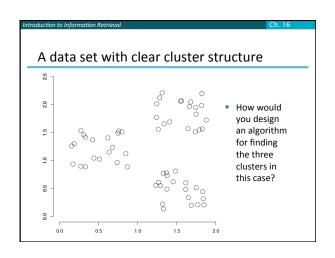
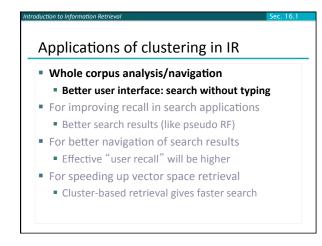
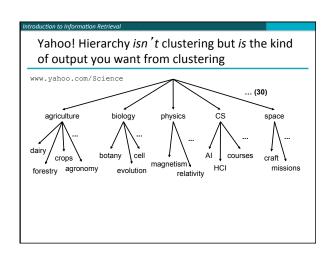
# Introduction to Information Retrieval Clustering Chris Manning and Pandu Nayak

# Today's Topic: Clustering Document clustering Motivations Document representations Success criteria Clustering algorithms Partitional Hierarchical

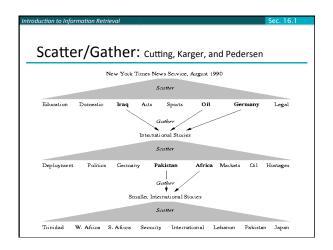


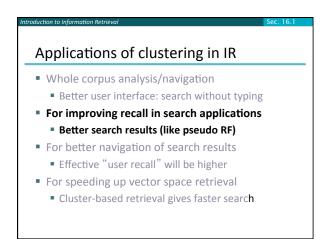


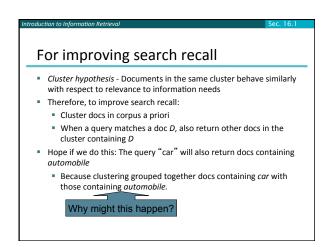


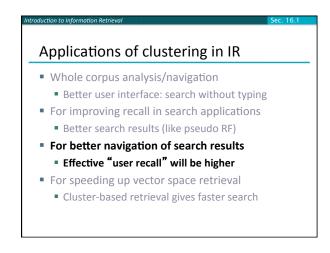


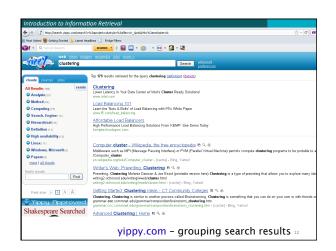












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- Applications of clustering in IR

  Whole corpus analysis/navigation
  - Better user interface: search without typing
- For improving recall in search applications
  - Better search results (like pseudo RF)
- For better navigation of search results
  - Effective "user recall" will be higher
- For speeding up vector space retrieval
  - Cluster-based retrieval gives faster search

Sec. 16.2

# Issues for clustering

- Representation for clustering
  - Document representation
    - Vector space? Normalization?
  - Need a notion of similarity/distance
- How many clusters?
  - Fixed a priori?
  - Completely data driven?
    - Avoid "trivial" clusters too large or small
      - If a cluster's too large, then for navigation purposes you've wasted an extra user click without whittling down the set of documents much.

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# Notion of similarity/distance

- Ideal: semantic similarity.
- Practical: term-statistical similarity (docs as vectors)
  - Cosine similarity
  - For many algorithms, easier to think in terms of a distance (rather than similarity) between docs.
  - We will mostly speak of Euclidean distance
    - But real implementations use cosine similarity

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## Hard vs. soft clustering

- Hard clustering: Each document belongs to exactly one cluster
  - More common and easier to do
- Soft clustering: A document can belong to more than one cluster.
  - Makes more sense for applications like creating browsable hierarchies
  - You may want to put a pair of sneakers in two clusters: (i) sports apparel and (ii) shoes
  - You can only do that with a soft clustering approach.
- We won't do soft clustering today. See IIR 16.5, 18

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# **Clustering Algorithms**

- Flat algorithms
  - Usually start with a random (partial) partitioning
  - Refine it iteratively
    - K means clustering
    - (Model based clustering)
- Hierarchical algorithms
  - Bottom-up, agglomerative
  - (Top-down, divisive)

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## **Partitioning Algorithms**

- Partitioning method: Construct a partition of n documents into a set of K clusters
- Given: a set of documents and the number K
- Find: a partition of *K* clusters that optimizes the chosen partitioning criterion
  - Globally optimal
    - Intractable for many objective functions
    - Ergo, exhaustively enumerate all partitions
  - Effective heuristic methods: K-means and K-medoids algorithms

See also Kleinberg NIPS 2002 – impossibility for natural clustering

# *K*-Means

- Assumes documents are real-valued vectors.
- Clusters based on centroids (aka the center of gravity or mean) of points in a cluster, c:

$$\vec{\mu}(c) = \frac{1}{\mid c \mid} \sum_{x \in c} \vec{x}$$

- Reassignment of instances to clusters is based on distance to the current cluster centroids.
  - (Or one can equivalently phrase it in terms of similarities)

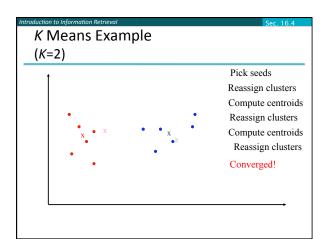


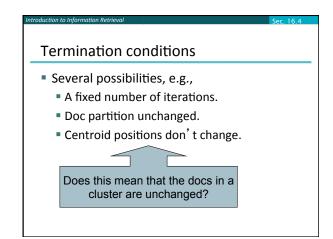
Select K random docs  $\{s_1, s_2, \dots s_K\}$  as seeds.

Until clustering *converges* (or other stopping criterion): For each doc  $d_i$ :

Assign  $d_i$  to the cluster  $c_j$  such that  $dist(x_i, s_j)$  is minimal. (Next, update the seeds to the centroid of each cluster) For each cluster  $c_i$ 

$$s_i = \mu(c_i)$$





# Convergence

- Why should the K-means algorithm ever reach a fixed point?
  - A state in which clusters don't change.
- K-means is a special case of a general procedure known as the Expectation Maximization (EM) algorithm.
  - EM is known to converge.
  - Number of iterations could be large.
    - But in practice usually isn't

# Convergence of K-Means

- Residual Sum of Squares (RSS), a goodness measure of a cluster, is the sum of squared distances from the cluster centroid:
  - RSS<sub>i</sub> =  $\Sigma_i ||d_i c_i||^2$  (sum over all  $d_i$  in cluster j)
- RSS =  $\Sigma_i$  RSS<sub>i</sub>
- Reassignment monotonically decreases RSS since each vector is assigned to the closest centroid.
- Recomputation also monotonically decreases each RSS<sub>j</sub> because ...

# Cluster recomputation in K-means

# Cluster recomputation in K-means

- RSS<sub>j</sub> =  $\Sigma_i \mid |d_i c_j| \mid^2 = \Sigma_i \Sigma_k (d_{ik} c_{jk})^2$ 
  - *i* ranges over documents in cluster *j*
- RSS<sub>j</sub> reaches minimum when:  $\Sigma_i - 2(d_{ik} - c_{ik}) = 0$  (for each  $c_{ik}$ )
- $m_i c_{ik} = \Sigma_i d_{ik}$  ( $m_i$  is # of docs in cluster j)
- $c_{ik} = (1/m_i) \Sigma_i d_{ik}$
- K-means typically converges quickly

# Time Complexity

- Computing distance between two docs is O(M) where M is the dimensionality of the vectors.
- Reassigning clusters: O(KN) distance computations, or O(KNM).
- Computing centroids: Each doc gets added once to some centroid: O(NM).
- Assume these two steps are each done once for I iterations: O(IKNM).

### Seed Choice Results can vary based on Example showing random seed selection. sensitivity to seeds Some seeds can result in poor convergence rate, or 0 convergence to sub-optimal In the above, if you start clusterings. with B and E as centroids you converge to {A,B,C} and {D,E,F} Select good seeds using a heuristic (e.g., doc least similar to any If you start with D and F existing mean) ou converge to Try out multiple starting points {A,B,D,E} {C,F} Initialize with the results of another method.

# K-means issues, variations, etc.

- Recomputing the centroid after every assignment (rather than after all points are re-assigned) can improve speed of convergence of K-means
- Assumes clusters are spherical in vector space
- Sensitive to coordinate changes, weighting etc.
- Disjoint and exhaustive
  - Doesn't have a notion of "outliers" by default
  - But can add outlier filtering

Dhillon et al. ICDM 2002 – variation to fix some issues with small

# How Many Clusters?

- Number of clusters K is given
  - lacktriangle Partition n docs into predetermined number of clusters
- Finding the "right" number of clusters is part of the problem
  - Given docs, partition into an "appropriate" number of subsets.
  - E.g., for query results ideal value of *K* not known up front though UI may impose limits.

# K not specified in advance

## 6 11 11 6

- Say, the results of a query.
- Solve an optimization problem: penalize having lots of clusters
  - application dependent, e.g., compressed summary of search results list.
- Tradeoff between having more clusters (better focus within each cluster) and having too many clusters

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## K not specified in advance

- Given a clustering, define the <u>Benefit</u> for a doc to be the cosine similarity to its centroid
- Define the <u>Total Benefit</u> to be the sum of the individual doc Benefits.

Why is there always a clustering of Total Benefit n?

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## Penalize lots of clusters

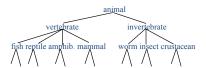
- For each cluster, we have a Cost C.
- Thus for a clustering with K clusters, the <u>Total Cost</u> is
- Define the <u>Value</u> of a clustering to be = Total Benefit - Total Cost.
- Find the clustering of highest value, over all choices of K.
  - Total benefit increases with increasing K. But can stop when it doesn't increase by "much". The Cost term enforces this.

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# Hierarchical Clustering

 Build a tree-based hierarchical taxonomy (dendrogram) from a set of documents.



 One approach: recursive application of a partitional clustering algorithm.

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## Dendrogram: Hierarchical Clustering

 Clustering obtained by cutting the dendrogram at a desired level: each connected component forms a cluster.



# Hierarchical Agglomerative Clustering

- Starts with each doc in a separate cluster
  - then repeatedly joins the <u>closest pair</u> of clusters, until there is only one cluster.
- The history of merging forms a binary tree or hierarchy.

Note: the resulting clusters are still "hard" and induce a partition

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## Closest pair of clusters

- Many variants to defining closest pair of clusters
- Single-link
  - Similarity of the most cosine-similar (single-link)
- Complete-link
  - Similarity of the "furthest" points, the *least* cosine-similar
- Centroid
  - Clusters whose centroids (centers of gravity) are the most cosine-similar
- Group average
  - Average cosine between all pairs of elements

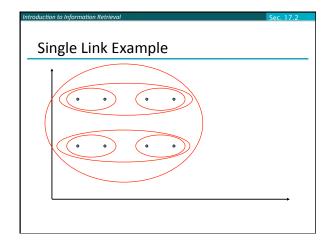
# Single Link Agglomerative Clustering

Use maximum similarity of pairs:

$$sim(c_i, c_j) = \max_{i \in S} sim(x, y)$$

- Can result in "straggly" (long and thin) clusters due to chaining effect.
- After merging  $c_i$  and  $c_j$ , the similarity of the resulting cluster to another cluster,  $c_k$ , is:

$$sim((c_i \cup c_j), c_k) = \max(sim(c_i, c_k), sim(c_j, c_k))$$



# Complete Link

Use minimum similarity of pairs:

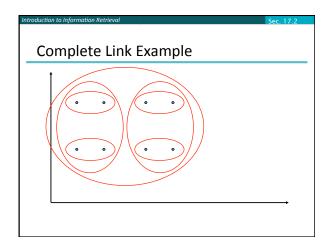
$$sim(c_i, c_j) = \min_{\substack{x \subseteq c_i, y \subseteq c_j}} sim(x, y)$$

• Makes "tighter," spherical clusters that are typically

- preferable.
- After merging  $c_i$  and  $c_i$ , the similarity of the resulting cluster to another cluster,  $c_k$ , is:

$$sim((c_i \cup c_j), c_k) = min(sim(c_i, c_k), sim(c_j, c_k))$$





# General HAC algorithm and complexity

1. Compute similarity between all pairs of documents

2. Do N – 1 times

 $O(N^2)$ 

1. Find closest pair of documents/clusters to merge

Naïve: O(N2) Priority Queue: O(N) Single link: O(N)

2. Update similarity of all documents/clusters to new cluster

persistent!

Naïve: O(N)

Priority Queue: Single link: O(N log N)

Best merge

# **Group Average**

Similarity of two clusters = average similarity of all pairs within merged cluster.

 $\frac{|c_i \cup c_j| (|c_i \cup c_j| - 1)}{|c_i \cup c_j|} \sum_{\vec{x} \in (c_i \cup c_j)} \sum_{\vec{y} \in (c_i \cup c_j); \vec{y} \neq \vec{x}} sim(\vec{x}, \vec{y})$ 

- Compromise between single and complete link.
- Two options:
  - Averaged across all ordered pairs in the merged cluster
  - Averaged over all pairs between the two original clusters
- No clear difference in efficacy

# Computing Group Average Similarity

Always maintain sum of vectors in each cluster.

$$\vec{s}(c_j) = \sum_{i=1}^{n} \vec{x}$$

 $\vec{s}(c_j) = \sum_{\vec{x} \in c_j} \vec{x}$  • Compute similarity of clusters in constant time:

$$sim(c_{i},c_{j}) = \frac{(\vec{s}(c_{i}) + \vec{s}(c_{j})) \bullet (\vec{s}(c_{i}) + \vec{s}(c_{j})) - (|c_{i}| + |c_{j}|)}{(|c_{i}| + |c_{j}|)(|c_{i}| + |c_{j}| - 1)}$$

# What Is A Good Clustering?

- Internal criterion: A good clustering will produce high quality clusters in which:
  - the intra-class (that is, intra-cluster) similarity is high
  - the inter-class similarity is low
  - The measured quality of a clustering depends on both the document representation and the similarity measure used

## External criteria for clustering quality

- Quality measured by its ability to discover some or all of the hidden patterns or latent classes in gold standard data
- Assesses a clustering with respect to ground truth ... requires labeled data
- Assume documents with C gold standard classes, while our clustering algorithms produce K clusters,  $\omega_1$ ,  $\omega_2$ , ...,  $\omega_K$  with  $n_i$  members.

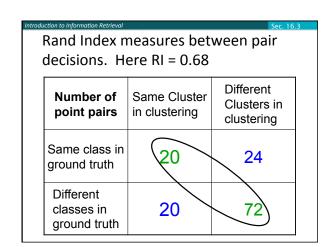
## **External Evaluation of Cluster Quality**

Simple measure: purity, the ratio between the dominant class in the cluster  $\omega_{i}$  and the size of cluster ω<sub>i</sub>

Purity 
$$(\omega_i) = \frac{1}{n_i} \max_j (n_{ij}) \quad j \in C$$

- Biased because having n clusters maximizes purity
- Others are entropy of classes in clusters (or mutual information between classes and clusters)

# Purity example Cluster III Cluster I Cluster II Cluster I: Purity = 1/6 (max(5, 1, 0)) = 5/6Cluster II: Purity = 1/6 (max(1, 4, 1)) = 4/6Cluster III: Purity = 1/5 (max(2, 0, 3)) = 3/5



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Rand index and Cluster F-measure

$$RI = \frac{A+D}{A+B+C+D}$$

Compare with standard Precision and Recall:

$$P = \frac{A}{A + B}$$

$$R = \frac{A}{A+C}$$

People also define and use a cluster F-measure, which is probably a better measure.

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## Final word and resources

- In clustering, clusters are inferred from the data without human input (unsupervised learning)
- However, in practice, it's a bit less clear: there are many ways of influencing the outcome of clustering: number of clusters, similarity measure, representation of documents, . . .
- Resources
  - IIR 16 except 16.5
  - IIR 17.1–17.3