# Unsupervised Learning and Data Mining

# Unsupervised Learning and Data Mining

Clustering

- Decision trees
- Artificial neural nets
- K-nearest neighbor
- Support vectors
- Linear regression
- Logistic regression

• ...

- F(x): true function (usually not known)
- D: training sample drawn from F(x)

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- F(x): true function (usually not known)
- D: training sample drawn from F(x)

G(x): model learned from training sample D

• Goal:  $E < (F(x)-G(x))^2 > is small (near zero) for future samples drawn from <math>F(x)$ 

Well Defined Goal:

Learn G(x) that is a good approximation to F(x) from training sample D

**Know How to Measure Error:** 

Accuracy, RMSE, ROC, Cross Entropy, ...

# Clustering



Supervised Learning

# Clustering

Unsupervised Learning

#### Train Set:

### Test Set:

### Train Set:

### Test Set:

### Train Set:

• • •

### Test Set

### Data Set:

• • •

# Supervised vs. Unsupervised Learning

### Supervised

- y=F(x): true function
- D: labeled training set
- D:  $\{x_i, y_i\}$
- y=G(x): model trained to predict labels D
- Goal:

$$E < (F(x)-G(x))^2 > \approx 0$$

• Well defined criteria: Accuracy, RMSE, ...

### Unsupervised

- Generator: true model
- D: unlabeled data sample
- D:  $\{x_i\}$
- Learn

??????????

• Goal:

??????????

• Well defined criteria:

### What to Learn/Discover?

- Statistical Summaries
- Generators
- Density Estimation
- Patterns/Rules
- Associations
- Clusters/Groups
- Exceptions/Outliers
- Changes in Patterns Over Time or Location

# Goals and Performance Criteria?

- Statistical Summaries
- Generators
- Density Estimation
- Patterns/Rules
- Associations
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# Clustering

# Clustering

- Given:
  - Data Set D (training set)
  - Similarity/distance metric/information
- Find:
  - Partitioning of data
  - Groups of similar/close items

# Similarity?

- Groups of similar customers
  - Similar demographics
  - Similar buying behavior
  - Similar health
- Similar products
  - Similar cost
  - Similar function
  - Similar store

**—** ...

• Similarity usually is domain/problem specific

# Types of Clustering

- Partitioning
  - K-means clustering
  - K-medoids clustering
  - EM (expectation maximization) clustering
- Hierarchical
  - Divisive clustering (top down)
  - Agglomerative clustering (bottom up)
- Density-Based Methods
  - Regions of dense points separated by sparser regions of relatively low density

# Types of Clustering

- Hard Clustering:
  - Each object is in one and only one cluster
- Soft Clustering:
  - Each object has a probability of being in each cluster

# Two Types of Data/Distance Info

• N-dim vector space representation and distance metric

...

```
Distance (D1,D2) = ???
```

- Pairwise distances between points (no N-dim space)
  - + Similarity/dissimilarity matrix (upper or lower diagonal)

```
+ Distance: 0 = \text{near}, \quad \infty = \text{far}
```

+ Similarity: 0 = far,  $\infty = near$ 

```
-- 1 2345678910
1 - ddddddddd
2 - dddddddd
3 - ddddddd
4 - ddddd
5 - ddddd
6 - dddd
7 - ddd
8 - dd
9 - d
```

# Agglomerative Clustering

• Put each item in its own cluster (641 singletons)

- Find all pairwise distances between clusters
- Merge the two *closest* clusters
- Repeat until everything is in one cluster

- Hierarchical clustering
- Yields a clustering with each possible # of clusters
- Greedy clustering: not optimal for any cluster size

# Agglomerative Clustering of Proteins



# Merging: Closest Clusters

- Nearest centroids
- Nearest medoids
- Nearest neighbors (shortest link)
- Nearest average distance (average link)
- Smallest greatest distance (maximum link)
- Domain specific similarity measure
  - word frequency, TFIDF, KL-divergence, ...
- Merge clusters that optimize criterion after merge
  - minimum mean\_point\_happiness

## Mean Distance Between Clusters

$$Mean\_Dist\ (c_1,c_2) = \frac{\displaystyle\sum_{i \in c_1} \sum_{j \in c_2} Dist\ (i,j)}{\displaystyle\sum_{i \in c_1} \sum_{j \in c_2} 1}$$

## Minimum Distance Between Clusters

$$Min\_Dist\ (c_1, c_2) = MIN_{i \in c_1, j \in c_2}(Dist\ (i, j))$$

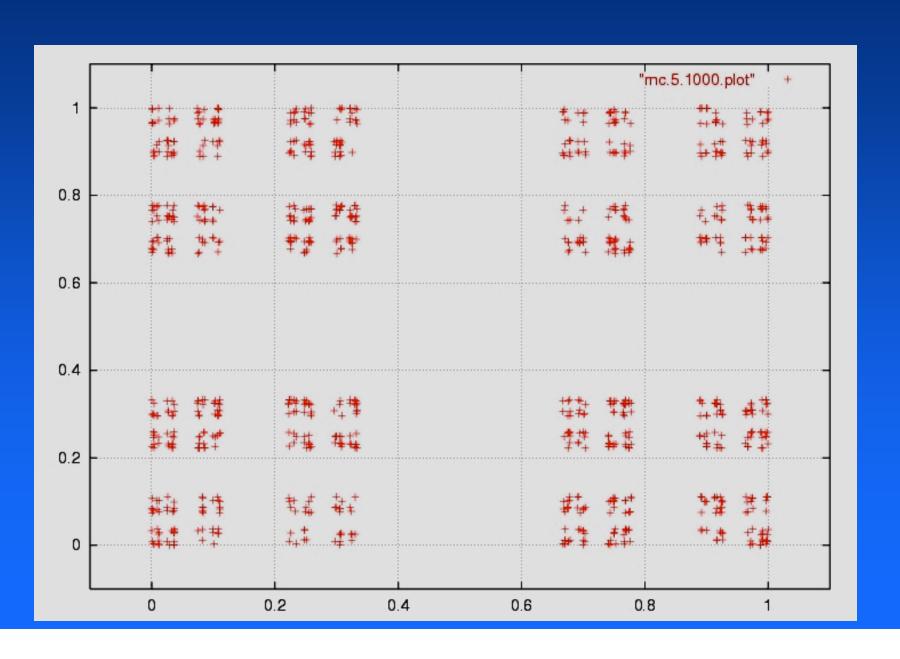
# Mean Internal Distance in Cluster

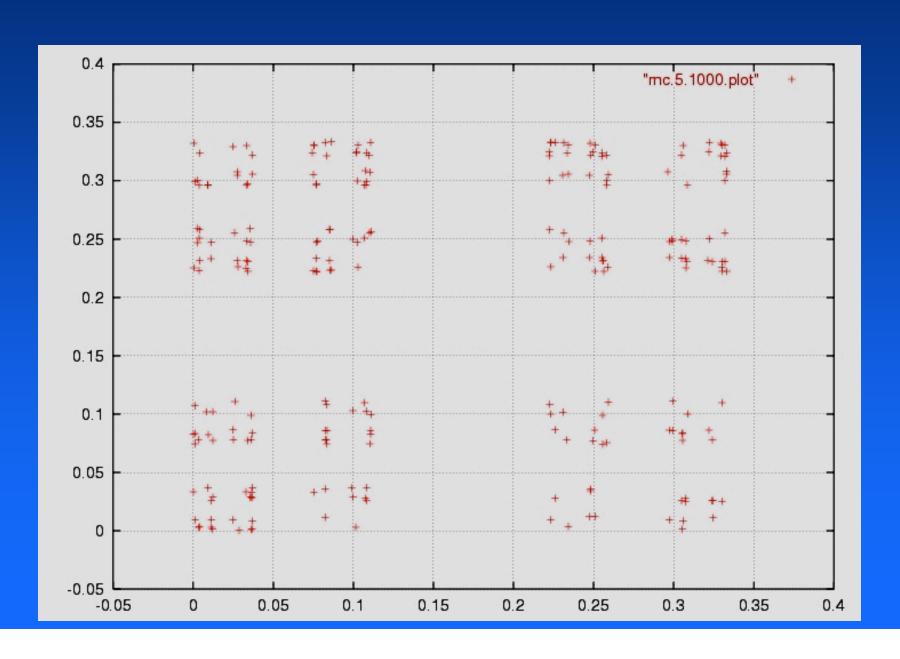
$$Mean\_Internal\_Dist(c) = \frac{\sum_{i \in c} \sum_{j \in c, i \neq j} Dist(i, j)}{\sum_{i \in c} \sum_{j \in c, i \neq j} 1}$$

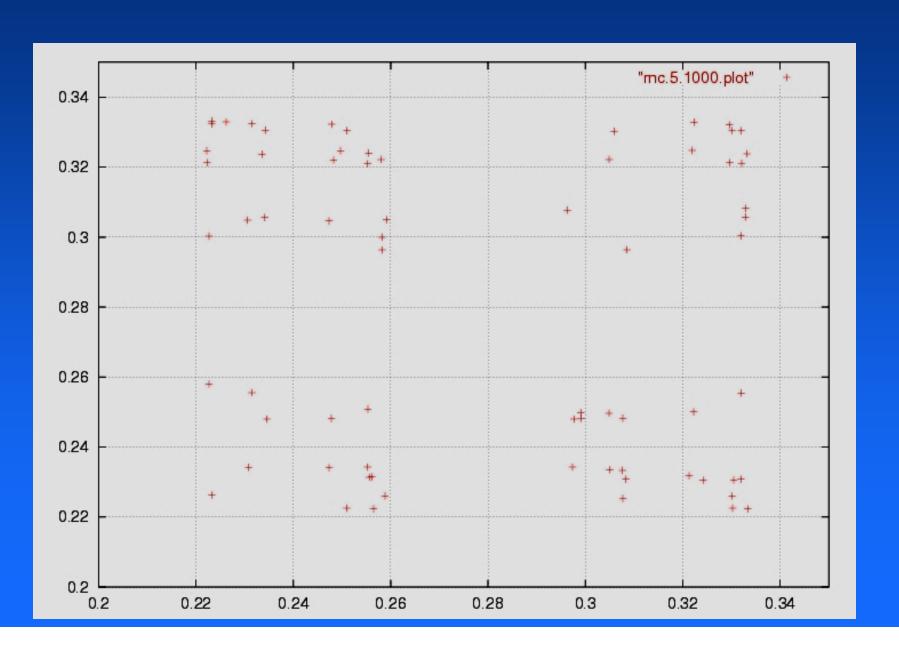
# Mean Point Happiness

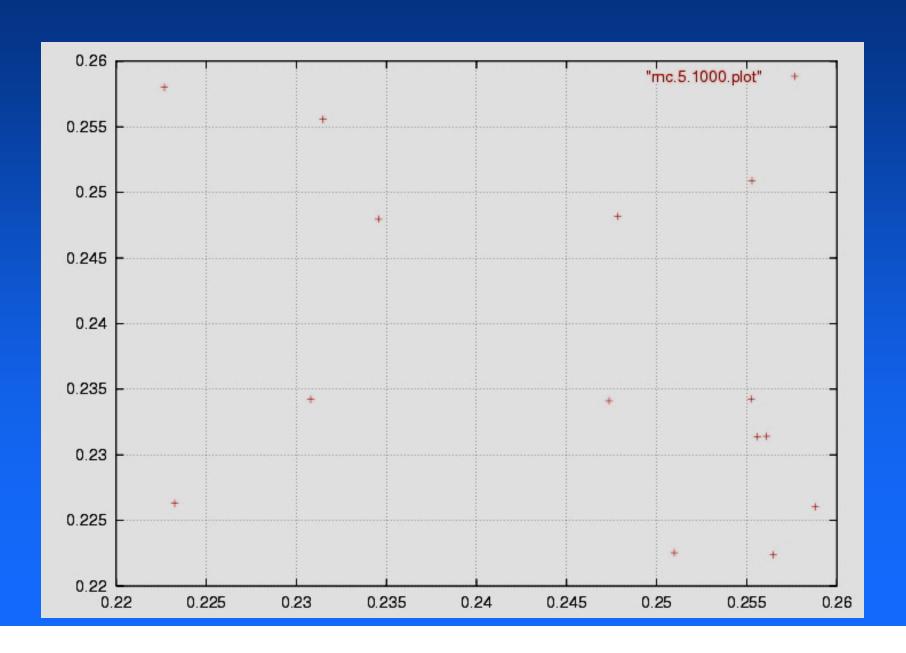
$$\delta_{ij} = \begin{cases} 1 & when \ cluster(i) = cluster(j) \\ 0 & when \ cluster(i) \neq cluster(j) \end{cases}$$

$$Mean\_Happiness = \frac{\sum_{i} \sum_{j} \delta_{ij} \quad Dist(i, j)}{\sum_{i} \sum_{j} \delta_{ij}}$$

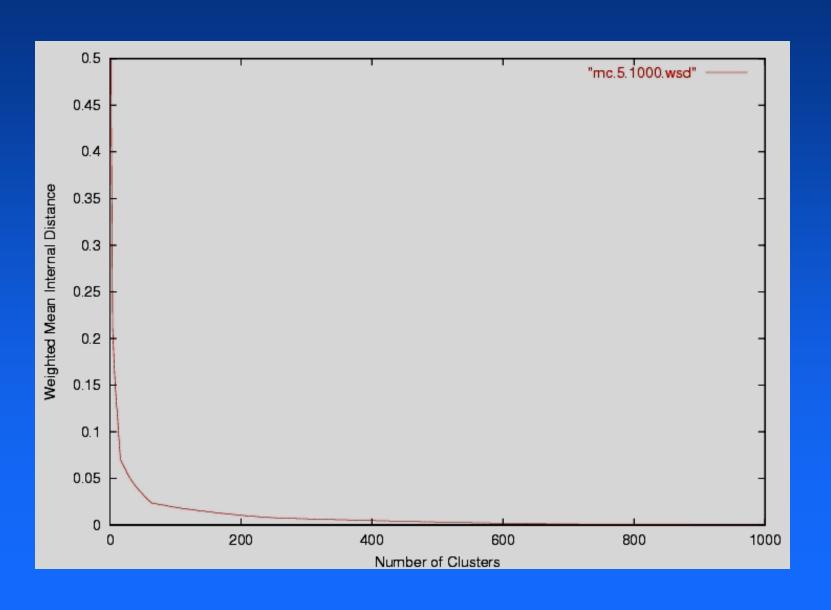




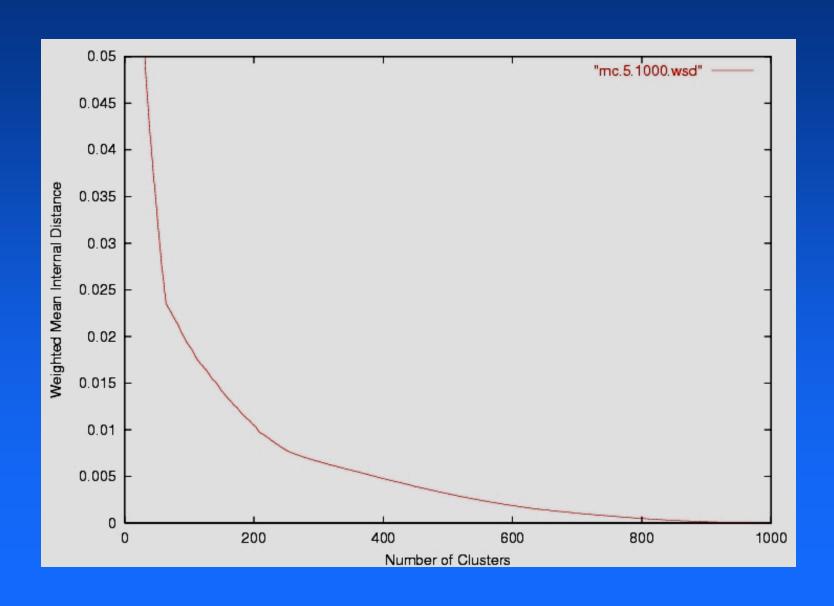




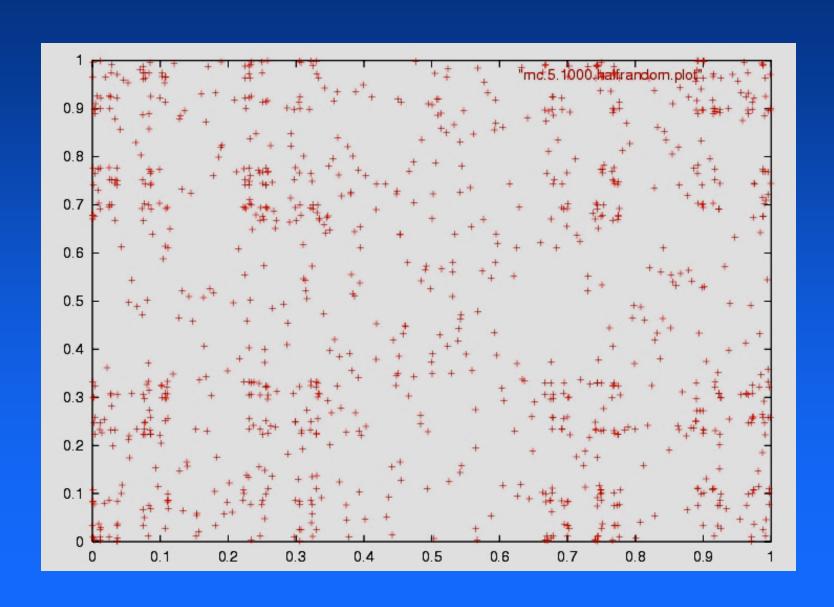
# Mean Point Happiness



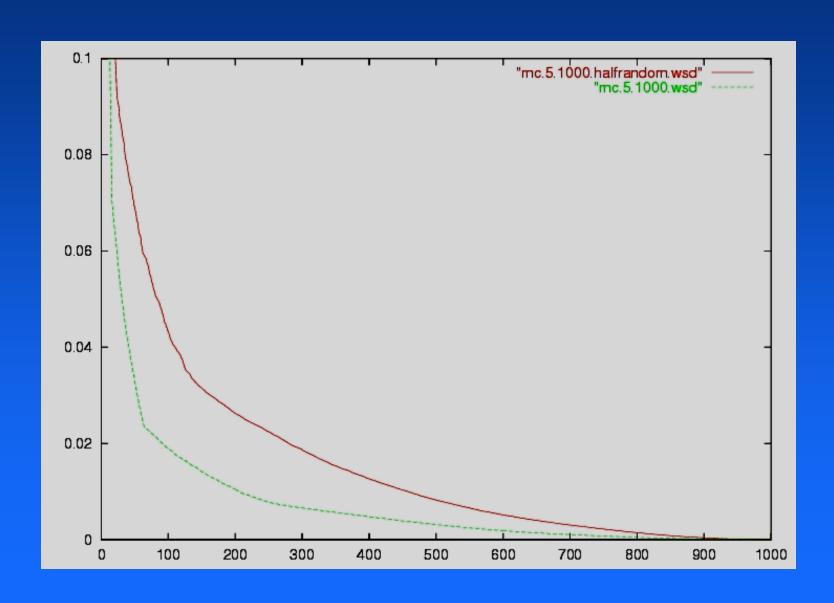
# Mean Point Happiness



# Recursive Clusters + Random Noise

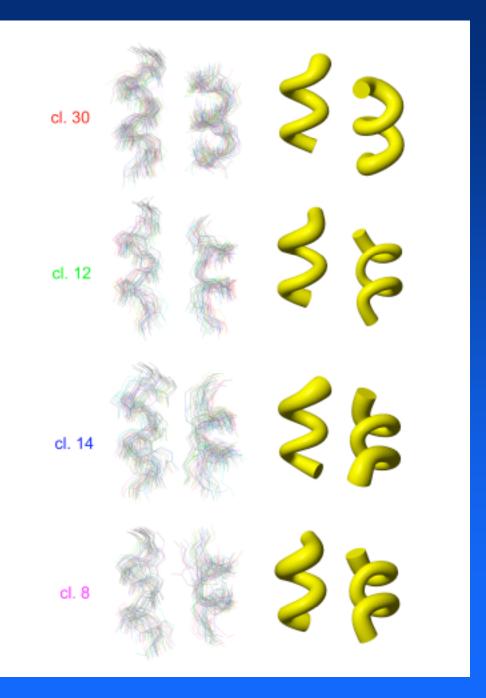


# Recursive Clusters + Random Noise



## Clustering Proteins





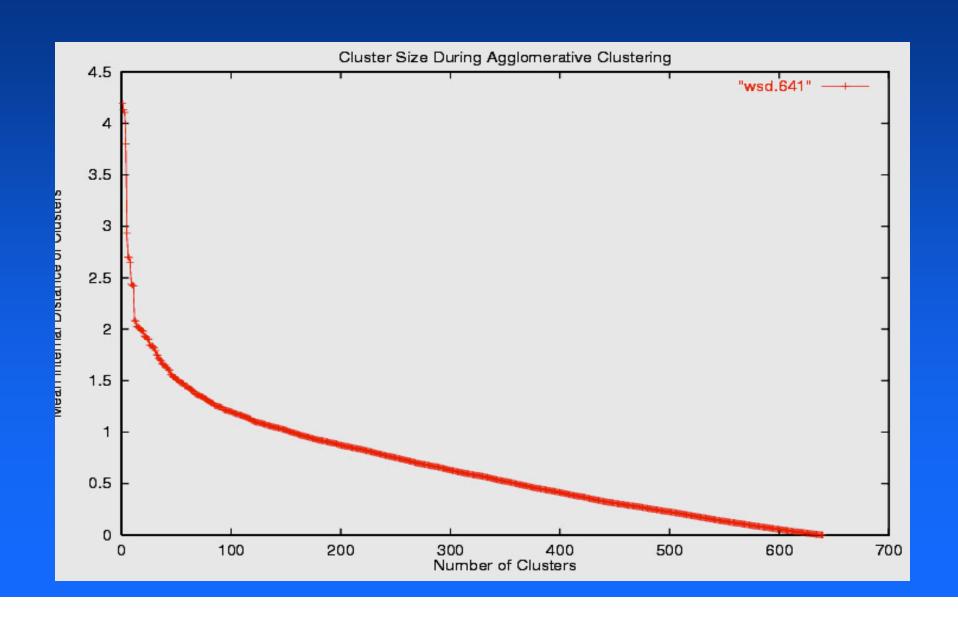
#### Distance Between Helices

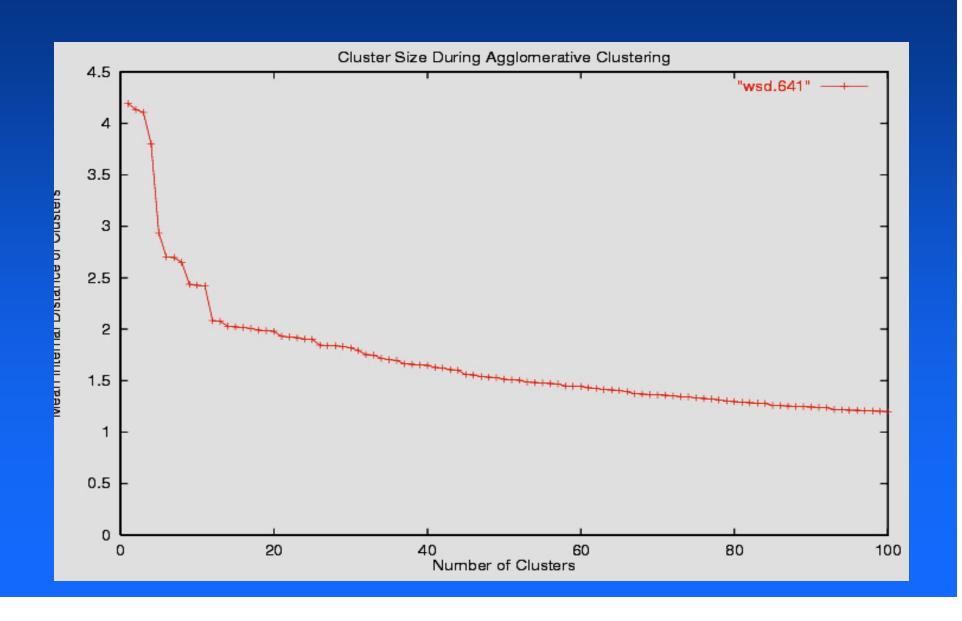
• Vector representation of protein data in 3-D space that gives x,y,z coordinates of each atom in helix

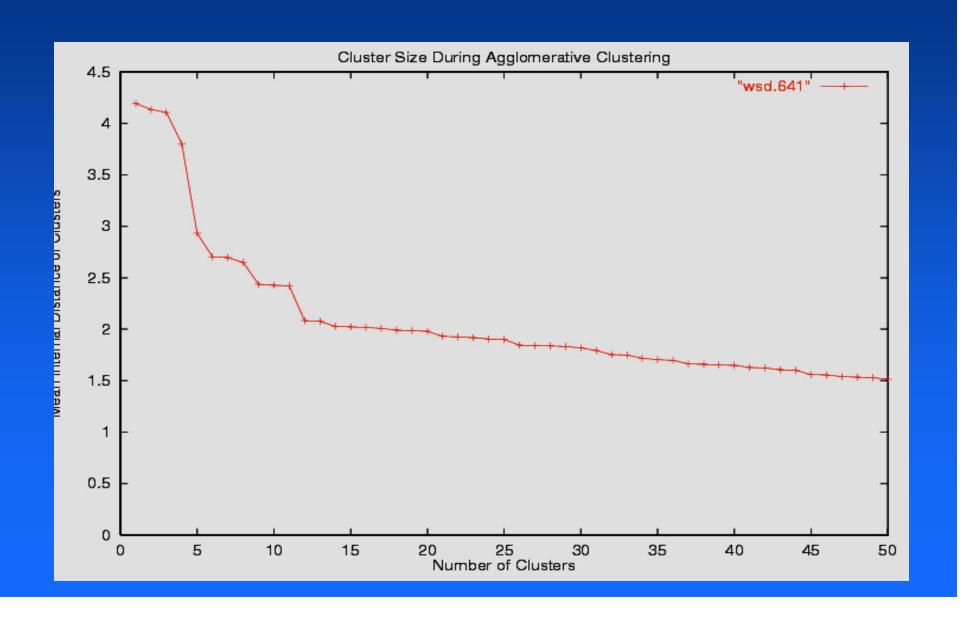
• Use a program developed by chemists (fortran) to convert 3-D atom coordinates into average atomic distances in angstroms between aligned helices

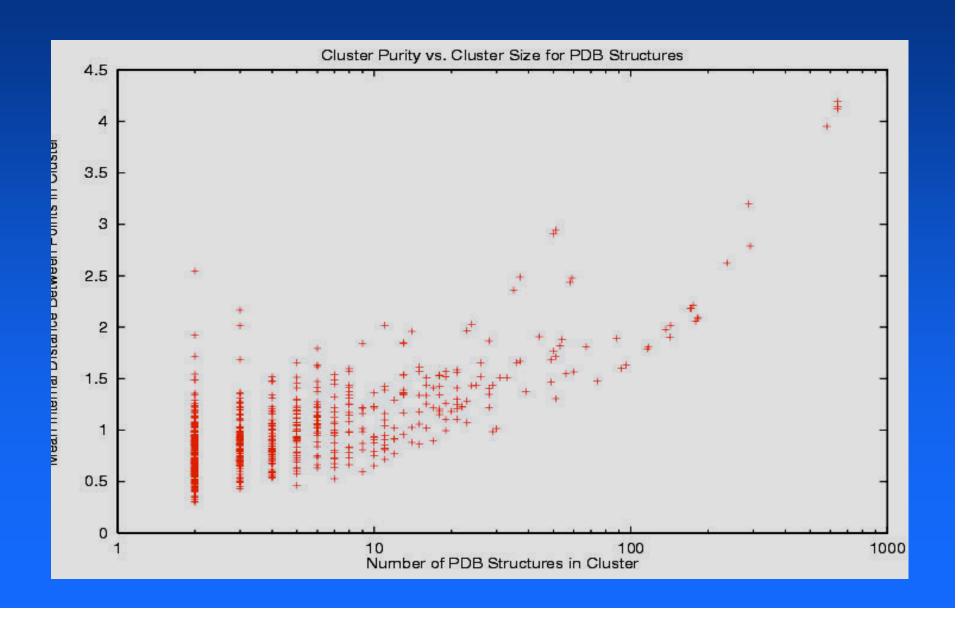
• 641 helices = 641 \* 640 / 2 = 205,120 pairwise distances



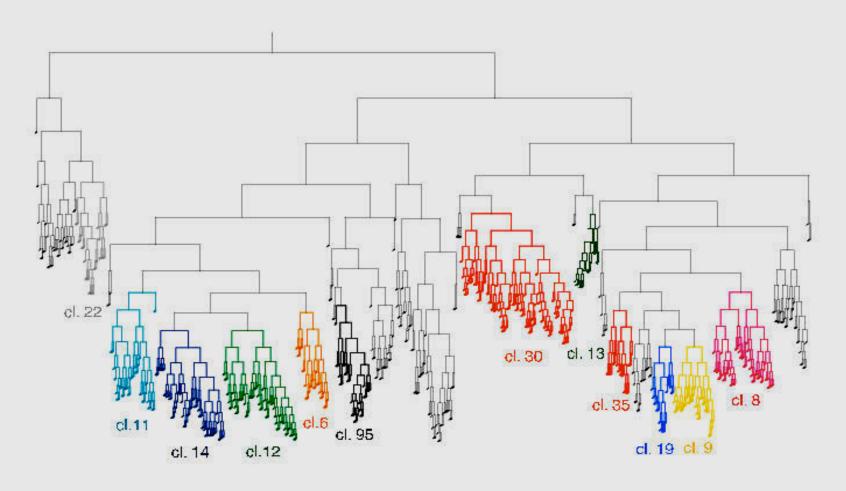




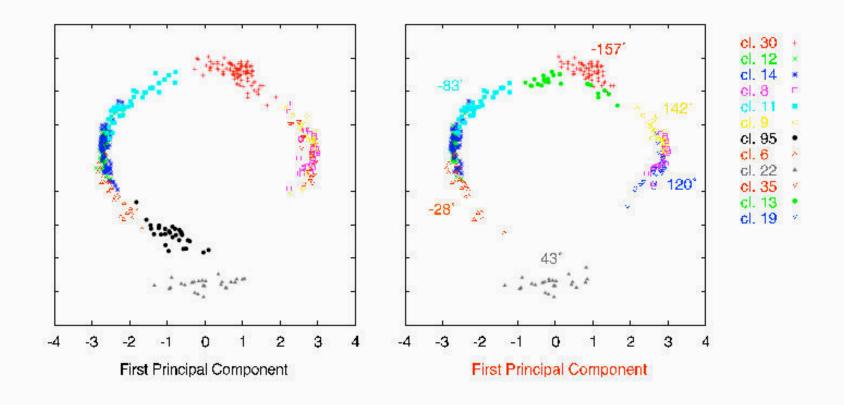




#### Agglomerative Clustering of Helix Pairs



#### Multidimensional Scaling of helix pairs by RMSD



#### Agglomerative Clustering

- Greedy clustering
  - once points are merged, never separated
  - suboptimal w.r.t. clustering criterion
- Combine greedy with iterative refinement
  - post processing
  - interleaved refinement

#### Agglomerative Clustering

- Computational Cost
  - O(N<sup>2</sup>) just to read/calculate pairwise distances
  - N-1 merges to build complete hierarchy
    - + scan pairwise distances to find closest
    - + calculate pairwise distances between clusters
    - + fewer clusters to scan as clusters get larger
  - Overall  $O(N^3)$  for simple implementations
- Improvements
  - sampling
  - dynamic sampling: add new points while merging
  - tricks for updating pairwise distances

#### K-Means Clustering

- Inputs: data set and k (number of clusters)
- Output: each point assigned to one of k clusters

- K-Means Algorithm:
  - -Initialize the k-means
    - +assign from randomly selected points
    - + randomly or equally distributed in space
  - -Assign each point to nearest mean
  - -Update means from assigned points
  - -Repeat until convergence

#### K-Means Clustering: Convergence

• Squared-Error Criterion

Squared 
$$\_Error = \sum_{c} \sum_{i \in c} (Dist(i, mean(c)))^{2}$$

- Converged when SE criterion stops changing
- Increasing K reduces SE can't determine K by finding minimum SE
- Instead, plot SE as function of K

#### K-Means Clustering

- Efficient
  - $-K \ll N$ , so assigning points is  $O(K*N) \ll O(N^2)$
  - updating means can be done during assignment
  - usually # of iterations << N</p>
  - Overall O(N\*K\*iterations) closer to O(N) than  $O(N^2)$
- Gets stuck in local minima
  - Sensitive to initialization
- Number of clusters must be pre-specified
- Requires vector space date to calculate means

#### Soft K-Means Clustering

- Instance of EM (Expectation Maximization)
- Like K-Means, except each point is assigned to each cluster with a probability
- Cluster means updated using weighted average
- Generalizes to Standard\_Deviation/Covariance
- Works well if cluster models are known

#### Soft K-Means Clustering (EM)

- -Initialize model parameters:
  + means
  + std\_devs
  + ...
- -Assign points probabilistically to each cluster
- -Update cluster parameters from weighted points
- -Repeat until convergence to local minimum

# What do we do if we can't calculate cluster means?

```
-- 1 2345678910

1 - ddddddddd

2 - dddddddd

3 - ddddddd

4 - dddddd

5 - ddddd

6 - dddd

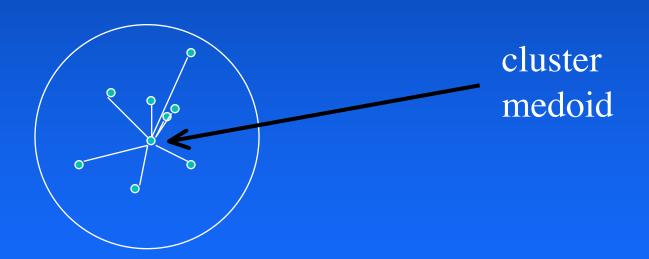
7 - dddd

8 - dd

9 - d
```

#### K-Medoids Clustering

$$Medoid(c) = pt \in c \ s.t. \ MIN(\sum_{i \in c} Dist(i, pt))$$



#### K-Medoids Clustering

- Inputs: data set and k (number of clusters)
- Output: each point assigned to one of k clusters

- Initialize k medoids
  - pick points randomly
- Pick medoid and non-medoid point at random
- Evaluate quality of swap
  - Mean point happiness
- Accept random swap if it improves cluster quality

#### Cost of K-Means Clustering

- n cases; d dimensions; k centers; i iterations
- compute distance each point to each center: O(n\*d\*k)
- assign each of n cases to closest center: O(n\*k)
- update centers (means) from assigned points: O(n\*d\*k)
- repeat i times until convergence
- overall: O(n\*d\*k\*i)
- much better than O(n<sup>2</sup>)-O(n<sup>3</sup>) for HAC
- sensitive to initialization run many times
- usually don't know k run many times with different k
- requires many passes through data set

# Graph-Based Clustering

#### Scaling Clustering to Big Databases

- K-means is still expensive: O(n\*d\*k\*I)
- Requires multiple passes through database
- Multiple scans may not be practical when:
  - database doesn't fit in memory
  - database is very large:
    - $+10^4$ - $10^9$  (or more) records
    - $+>10^2$  attributes
  - expensive join over distributed databases

#### Goals

- 1 scan of database
- early termination, on-line, anytime algorithm yields current best answer

#### Scale-Up Clustering?

- Large number of cases (big n)
- Large number of attributes (big d)
- Large number of clusters (big c)