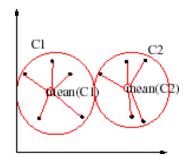
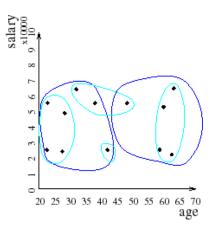
K-means and K-medoids algorithms

- Minimizes the sum of square distances of points to cluster representative
- Efficient iterative algorithms (O(n))



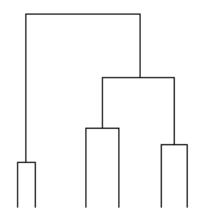
Problems with K-means type algorithms

- Clusters are approximately spherical
- High dimensionality is a problem
- The value of K is an input parameter



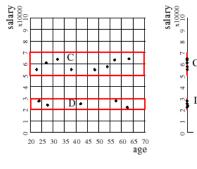
Hierarchical Clustering

- Quadratic algorithms
- Running time can be improved using sampling [Guha et al, SIGMOD 1998] [Kollios et al, ICDE 2001]



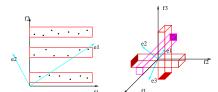
Density Based Algorithms

- Clusters are regions of space which have a high density of points
- Clusters can have arbitrary shapes



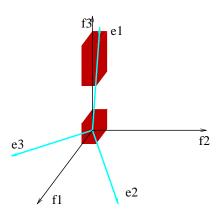
Dimensionality Reduction

- Reduce the dimensionality of the space, while preserving distances
- Many techniques (SVD, MDS)
- May or may not help



Dimensionality Reduction

• Example: SVD decomposition



Speeding up the clustering algorithms: Data Reduction

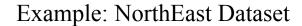
- Data Reduction:
 - approximate the original dataset using a small representation
 - ideally, the representation must be stored in main memory
 - summarization, compression
- The accuracy loss must be as small as possible.
- Use the approximated dataset to run the clustering algorithms

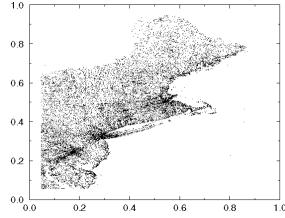
Random Sampling as a Data Reduction Method

- Random Sampling is used as a data reduction method
- Idea: Use a random sample of the dataset and run the clustering algorithm over the sample
- Used for clustering and association rule detection [Ng and Han 94][Toivonen 96][Guha et al 98]
- But:
 - For datasets that contain clusters with different densities, we may miss some sparse ones
 - For datasets with noise we may include significant amount of noise in our sample

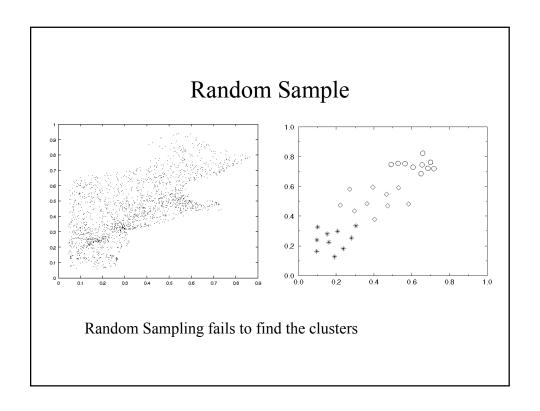
A better idea: Biased Sampling

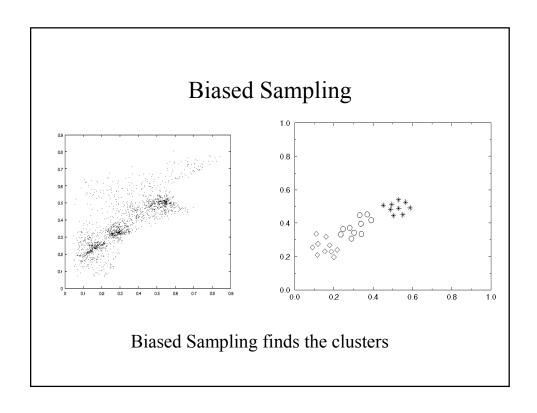
- Use biased sampling instead of random sampling
- In biased sampling, the prob that a point is included in the sample depends on the local density
- We can oversample or undersample regions in our datasets depending on the DM task at hand





NorthEast Dataset, 130K postal addresses in North Eastern USA





The Biased Sampling Technique

- Basic idea:
 - First compute an approximation of the density function of the dataset
 - Use the density function to define the bias for each point and perform the sampling

[Kollios et al, ICDE 2001]

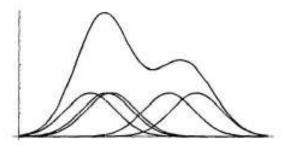
[Domeniconi and Gunopulos, ICML 2001]

[Palmer and Faloutsos, SIGMOD 2000]

Density Estimation

- We use kernels to approximate the probability density function (pdf)
- We scan the dataset and we compute an initial random sample and standard deviation
- For each sample we use a kernel. The approximate pdf is the sum of all kernels

Kernel Estimator



Example of a Kernel Estimator

The sampling step

- Let f(p) the pdf value for the point $p \in D$ $p = (x_1, x_2, ..., x_d)$
- We define $L(p) = f(p)^{\alpha}$, where α is parameter
- We compute the normalization parameter k (in one scan):

$$k = \sum_{p \in D} L(p)$$

The sampling step (cont.)

• The sampling bias is proportional to:

$$\frac{b}{k}L(\mathbf{p})$$

Where b is the size of the sample and k the normalization factor

- In another scan we perform the sampling (two scans)
- We can combine the above two steps into one scan

The variable α

- If $\alpha = 0$ then we have uniform random sampling bias: $\frac{b}{}$
- If $\alpha > 0$ then regions with higher density are sampled at a higher rate
- If α < 0 then regions with higher density are sampled at a lower rate

Bias
$$\sim \frac{b}{k} f(\mathbf{p})^a$$

• We can show that if $\alpha > -1$, relative densities are preserved in the sample

