## Database and Information Systems

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## Indexing Structures for High-Dimensional Data

#### Readings:

Schmitt, I.: Ähnlichkeitssuche in Multimedia-Datenbanken. Retrieval, Suchalgorithmen und Anfragebehandlung. Oldenbourg, München 2006.

### Indexing Structures for High-Dimensional Data

- Queries in High-Dimensional Databases
- Boundary-Based Indexes
- Dimensionality Reduction

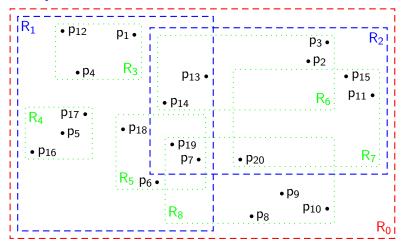
- range queries: find all objects whose attribute values fall within certain given ranges
  - rectangular hyper-window (window query)
- similarity range queries: find all objects which are within a given distance from an object
  - hyper-sphere query
  - distance is defined based on an application specific metrics
- nearest neighbor query: find the object which is closest to a given object
- reverse nearest neighbor query: find all objects for which a given object would be a nearest neighbor
  - nn-relation is not symmetric
  - e.g. finding an optimal location for a meeting

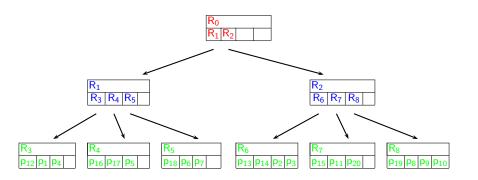
- k-nearest neighbor (KNN) queries: find the k-most similar objects which are closest in distance to a given object
  - high dimensional data have low contrast in distance
  - ullet if more similar objects than required exist ightarrow random choice
- similarity join: find all pairs of objects which are similar enough
  - distance is smaller than a predefined threshold
- similarity queries can be emulated by range queries using a filter-and-refine approach
  - filter: find the candidates with a sufficiently large bounding box
  - refine: check the similarity criterion for each of them

- requirements for index structures
  - soundness and completeness
  - suitability for high-dimensional problem spaces
  - suitability for spatially extended objects
  - retrieval efficiency
  - efficient update operations (insertion, deletion, update)
  - support for several distance metrics
  - minimal space requirements
- ullet no one-fits-all solution o compromise needs to be found

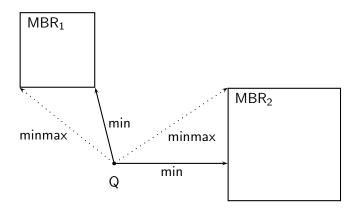
- curse of high dimensionality
  - a high-dimensional data space is sparse
  - distance between data points increases
  - data contrast is low
    - distance between the nearest and the farthest data point is reduced
    - high number of almost equally similar data points
  - notion of similarity vanishes
  - approximate (probabilistic) methods suffice

- range queries: R-Trees
  - multi-dimensional extension of B<sup>+</sup>-trees
  - preserves height balance
  - insertion: node splitting
  - deletion: node merging
- leaf nodes
  - object identifier
  - bounding box: Minimum Bounding Rectangle (MBR)
- non-leaf nodes
  - child-pointer
  - bounding box for the whole sub-tree





- range queries: recursive tree traversal
  - find all leaf nodes where the bounding box overlaps the query range
  - non-deterministic search: all non-leaf nodes with intersecting bounding boxes have to be considered
- nn-queries: objects in a node can be ordered
  - min-distance:
    minimal distance between a query point and a point in an MBR optimistic expectation for the distance to the nearest neighbor
  - minmax-distance:
     maximal distance a query point can have to a nearest neighbor
     in an MBR
     pessimistic expectation for the distance to the nearest neighbor



- search maintains an upper limit for the best result found so far
- $min(Q,MBR_1) > minmax(Q,MBR_2)$ 
  - MBR<sub>1</sub> need not be considered
- upper limit > minmax(Q,MBR<sub>1</sub>)
  - MBR<sub>1</sub> contains a closer neighbor
  - upper limit can be updated to minmaxdist(Q,MBR<sub>1</sub>)
- upper limit < min(Q,MBR<sub>1</sub>)
  - MBR<sub>1</sub> need not be considered
- can be extended to deal with knn-queries
  - maintaining the candidates in a priority queue of length k
  - upper limit refers to the last entry in the priority queue

- node insertion
  - least coverage criterion: choose the branch which requires minimal enlargement to also accommodate the new object
  - in case of a tie: choose the smallest box
  - all traversed and splitted nodes are readjusted to a minimum bounding box
- node splitting: like in a B-tree
- node deletion:
  - changes the bounding box in ancestor nodes
    - ightarrow adaptation needed
  - in case of underflow: delete the node and reinsert its remaining childs from the root
    - might cause further node deletions

- problems:
  - the overlap of bounding boxes increases as the dimensionality grows
    - → not well suited for multi-dimensional problems
    - $\rightarrow$  for a large number of dimensions (d > 10) a sequential scan can be shown to be more efficient
  - overlap is sensitive to the order of insertion → reorganisation can give an advantage (e.g. node re-insertion from the root)
  - not well suited for KNN queries
  - only for vector spaces with a EUCLIDEAN distance

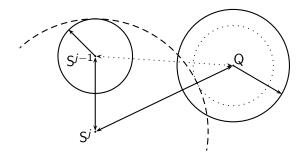
- variants
  - R<sup>+</sup>-Tree: no overlaps allowed  $\rightarrow$  too many splits
  - R\*-Tree: optimizes the margin of the bounding boxes squarish boxes are preferred
  - X-tree: avoiding splits if they result in highly overlapping nodes

     → supernodes
  - A-Trees: using virtual bounding boxes to approximate the minimal ones

- alternative: using bounding spheres instead of bounding boxes (SS-Tree)
  - · centroid: mean vector
  - radius: distance from the centroid to the farthest data point
- generalized version: metric tree (M-Tree)
  - data are clustered first
  - cluster are mapped to nodes in the tree
  - centroids are used as routing objects
  - triangle inequation is used to exclude subtrees from being searched
  - tree is balanced

- advantages
  - works with arbitraty metrics
  - lower dimensionality: d+1 instead of 2d
  - radius of the bounding sphere is determined by the distance
    - $\rightarrow$  insensitive to the dimensionality while diagonal in a box increases with dimensionality
      - but: boxes can be better adapted to different value ranges along different dimensions
  - · well suited for similarity search

 distance between a centroid and the query can be used to prune the search space



- there must be a subcluster in the closer hemisphere
- the true distance to the subcluster cannot be larger than the distance to the centroid it belongs to

- kd-Tree: binary search tree
- splitting on different levels of the tree can be done along different dimensions of the feature space
- partitions the feature space completely (no overlaps)
- tree is unbalanced (by definition)
- hyper-rectangles are usually larger than necessary but queried more often
  - ightarrow worse performance

- no tree-based index is efficient for truly high-dimensional problems
- common assumption:
  - data points can be clustered
  - certain clusters can be excluded from search
- assumption is fundamentally wrong for similarity search in a space with many uncorrelated features

### **Dimensionality Reduction**

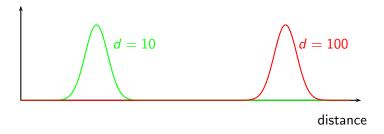
- dimensionality reduction techniques are always lossy
- indexing based on important attributes: TV-Tree (Telescopic-Vector Tree)
  - similar to an R-Tree, but nodes higher up in the tree use fewer features
  - features have to be selected according to some predefined ranking
- Principal Component Analysis
  - decorrelation of features
  - transforms the whole feature space
  - preserves similarity properties
  - using only a subset of the transformed feature space for indexing
  - efficient if data dimensions are globally correlated
  - but degree of correlation might change for dynamic data sets

## **Dimensionality Reduction**

- alternative: Local Dimensionality Reduction (LDR)
  - detect local clusters in the data set
  - perform a LDR for the individual clusters
  - build a local, low dimensional index using the transformed feature space
  - build a global index for the clusters
  - data points which do not belong to any cluster are treated as outliers and cannot be indexed at all
  - user can determine the amount of information loss, which affects the query precision and the query costs
- general problems:
  - degree of correlation might change for dynamic data sets
  - approach is also based on the idea of clustering

# The Curse of Dimensionality (Revisited)

- $\bullet$  for high dimensional problems in a  $\operatorname{EuclideAN}$  space the
  - expected value for the distance between two data points grows
  - but the standard deviation is constant



- many data points have a similar distance
- any kind of clustering becomes impossible

# The Curse of Dimensionality (Revisited)

- approximation error:
  - average distance between
    - a point query and the nearest data point of a cluster and
    - the point query and the cluster itself
- approximation error grows linearly with the expected value of the distance and
- eventually exeeds the greatest data point distance!
- → there is no method that reliably justifies to exclude a cluster from being considered in a high-dimensional data space

#### Signature-based Access

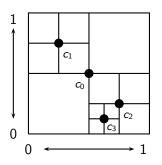
- vector approximation techniques (VA file)
- assumption: space can be partitioned into a finite set of discrete cells
  - partition the data space
    - number of cells per dimension is given
    - boundaries are chosen in a way that the number of data points in each intervall is evenly distributed
  - store the cell boundaries
  - represent each cell by a signature
    - i.e. concatenation of a binary representation for the cell number in each dimension
  - store the signature together with the full vector for each data point
  - use the signature as a filter to eliminate the data points that are not in the answer set

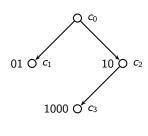
#### Signature-based Access

- search:
  - determine the relevant cells by a sequential scan
  - fetch the data points within the cells from the data base
  - check the search conditions for the fetched items
- but number of bits needs careful tuning
- drawback: fixed number of bits to describe a data point

## Signature-based Access

- modification: active vertice file (AV file)
  - partition the data space into a hierarchy of cells with different granualarity

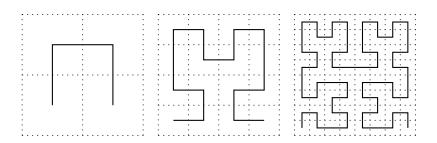




- a data point is assigned to a cell if it is closer than a given radius r from its centroid
- choice of *r* determines the efficiency

- special case of vector approximation
  - locality preserving total ordering of points in a k-dimensional space
  - e.g. Hilbert-curve
  - special enumeration scheme for the cells, e.g. for a 2-dimensional space:

- segment the space into 4  $(2^k)$  cells
- order the cells according to a basic curve
- recursively call the algorithm on all the cells until a detailed enough segmentation has been reached



• enumerate the cells along the path

	25 – 26 -   · · ·	· -   <del> </del>	· -   <del> </del>
19 – 18	· · · · · · · · · · · · · · · · · · ·	35 <del>-</del> 34	45 <del>-</del> 44
16 – 17			
14 - 13	8 <del></del> 9	54 <del>-</del> 55	50 <b>–</b> 49
1 — 2	7 — 6 - 4 — 5		61 – 62

- not all regions in the original space can be represented as contiguous regions in the new one
- → decomposition into several regions necessary

5 <b>-</b>		9 — 10
4	7 –	8 11
3 -	<del>-</del> 2 :	13 - 12
0 –		14 – 15

4 : 7 -	9 — 10 + 8 ] [11]
	13 - 12 13 - 12 14 - 15