

A more radical file organization is to regard the fact table as a large multidimensional array and store it and index it as such. This approach is taken in MOLAP systems. Since the array is much larger than available main memory, it is broken up into contiguous chunks, as discussed in Section 23.8. In addition, traditional B+ tree indexes are created to enable quick retrieval of chunks that contain tuples with values in a given range for one or more dimensions.

25.7 DATA WAREHOUSING

Data warehouses contain consolidated data from many sources, augmented with summary information and covering a long time period. Warehouses are much larger than other kinds of databases; sizes ranging from several gigabytes to terabytes are common. Typical workloads involve ad hoc, fairly complex queries and fast response times are important. These characteristics differentiate warehouse applications from OLTP applications, and different DBMS design and implementation techniques must be used to achieve satisfactory results. A distributed DBMS with good scalability and high availability (achieved by storing tables redundantly at more than one site) is required for very large warehouses.

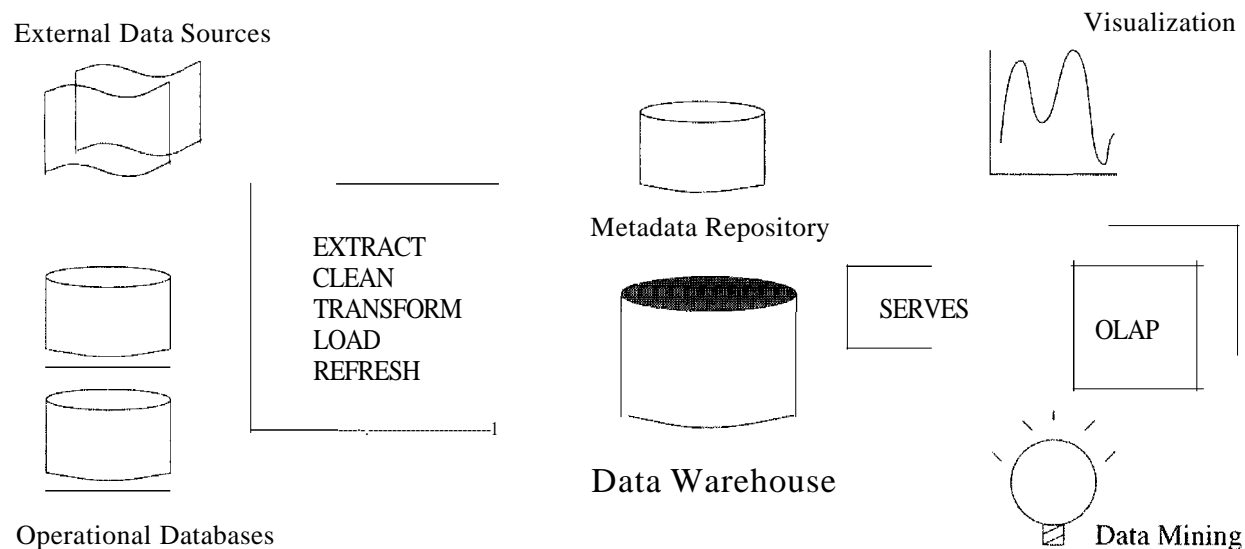


Figure 25.10 A typical Data Warehousing Architecture

A typical data warehousing architecture is illustrated in Figure 25.10. An organization's daily operations access and modify operational databases. Data from these operational databases and other external sources (e.g., customer profiles supplied by external consultants) are extracted by using interfaces such as JDBC (see Section 6.2).

25.7.1 Creating and Maintaining a Warehouse

Many challenges must be met in creating and maintaining a large data warehouse. A good database schema must be designed to hold an integrated collection of data copied from diverse sources. For example, a company warehouse might include the inventory and personnel departments' databases, together with sales databases maintained by offices in different countries. Since the source databases are often created and maintained by different groups, there are a number of semantic mismatches across these databases, such as different currency units, different names for the same attribute, and differences in how tables are normalized or structured; these differences must be reconciled when data is brought into the warehouse. After the warehouse schema is designed, the warehouse must be populated, and over time, it must be kept consistent with the source databases.

Data is extracted from operational databases and external sources, cleaned to minimize errors and fill in missing information when possible, and **transformed** to reconcile semantic mismatches. Transforming data is typically accomplished by defining a relational view over the tables in the data sources (the operational databases and other external sources). Loading data consists of materializing such views and storing them in the warehouse. Unlike a standard view in a relational DBMS, therefore, the view is stored in a database (the warehouse) that is different from the database(s) containing the tables it is defined over.

The cleaned and transformed data is finally loaded into the warehouse. Additional preprocessing such as sorting and generation of summary information is carried out at this stage. Data is partitioned and indexes are built for efficiency. Due to the large volume of data, loading is a slow process. Loading a terabyte of data sequentially can take weeks, and loading even a gigabyte can take hours. Parallelism is therefore important for loading warehouses.

After data is loaded into a warehouse, additional measures must be taken to ensure that the data in the warehouse is periodically refreshed to reflect updates to the data sources and periodically purge old data (perhaps onto archival media). Observe the connection between the problem of refreshing warehouse tables and asynchronously maintaining replicas of tables in a distributed DBMS. Maintaining replicas of source relations is an essential part of warehousing, and this application domain is an important factor in the popularity of asynchronous replication (Section 22.11.2), even though asynchronous replication violates the principle of distributed data independence. The problem of refreshing warehouse tables (which are materialized views over tables in

the source databases) has also renewed interest in incremental maintenance of materialized views. (We discuss materialized views in Section 25.8.)

An important task in maintaining a warehouse is keeping track of the data currently stored in it; this bookkeeping is done by storing information about the warehouse data in the system catalogs. The system catalogs associated with a warehouse are very large and often stored and managed in a separate database called a metadata repository. The size and complexity of the catalogs is in part due to the size and complexity of the warehouse itself and in part because a lot of administrative information must be maintained. For example, we must keep track of the source of each warehouse table and when it was last refreshed, in addition to describing its fields.

The value of a warehouse is ultimately in the analysis it enables. The data in a warehouse is typically accessed and analyzed using a variety of tools, including OLAP query engines, data mining algorithms, information visualization tools, statistical packages, and report generators.

25.8 VIEWS AND DECISION SUPPORT

Views are widely used in decision support applications. Different groups of analysts within an organization are typically concerned with different aspects of the business, and it is convenient to define views that give each group insight into the business details that concern it. Once a view is defined, we can write queries or new view definitions that use it, as we saw in Section 3.6; in this respect a view is just like a base table. Evaluating queries posed against views is very important for decision support applications. In this section, we consider how such queries can be evaluated efficiently after placing views within the context of decision support applications.

25.8.1 Views, OLAP, and Warehousing

Views are closely related to OLAP and data warehousing.

OLAP queries are typically aggregate queries. Analysts want fast answers to these queries over very large datasets, and it is natural to consider precomputing views (see Sections 25.9 and 25.10). In particular, the CUBE operator—discussed in Section 25.3—gives rise to several aggregate queries that are closely related. The relationships that exist between the many aggregate queries that arise from a single CUBE operation can be exploited to develop very effective precomputation strategies. The idea is to choose a subset of the aggregate queries for materialization in such a way that typical CUBE queries can be quickly answered by using the materialized views and doing some additional computation. The

choice of views to materialize is influenced by how many queries they can potentially speed up and by the amount of space required to store the materialized view (since we have to work with a given amount of storage space).

A data warehouse is just a collection of asynchronously replicated tables and periodically synchronized views. A warehouse is characterized by its size, the number of tables involved, and the fact that most of the underlying tables are from external, independently maintained databases. Nonetheless, the fundamental problem in warehouse maintenance is asynchronous maintenance of replicated tables and materialized views (see Section 25.10).

25.8.2 Queries over Views

Consider the following view, `RegionalSales`, which computes sales of products by category and state:

```
CREATE VIEW RegionalSales (category, sales, state)
AS SELECT P.category, S.sales, L.state
   FROM   Products P, Sales S, Locations L
   WHERE  P.pid = S.pid AND S.locid = L.locid
```

The following query computes the total sales for each category by state:

```
SELECT  R.category, R.state, SUM (R.sales)
FROM    RegionalSales R
GROUP BY R.category, R.state
```

While the SQL standard does not specify how to evaluate queries on views, it is useful to think in terms of a process called *query modification*. The idea is to replace the occurrence of `RegionalSales` in the query by the view definition. The result on this query is

```
SELECT  H.category, R.state, SUM (R.sales)
FROM    ( SELECT P.category, S.sales, L.state
          FROM   Products P, Sales S, Locations L
          WHERE  P.pid = S.pid AND S.locid = L.locid ) AS R
GROUP BY R.category, R.state
```

25.9 VIEW MATERIALIZATION

We can answer a query on a view by using the query modification technique just described. Often, however, queries against complex view definitions illustrate

be answered very fast because users engaged in decision support activities require interactive response times. Even with sophisticated optimization and evaluation techniques, there is a limit to how fast we can answer such queries. Also, if the underlying tables are in a remote database, the query modification approach may not even be feasible because of issues like connectivity and availability.

An alternative to query modification is to precompute the view definition and store the result. When a query is posed on the view, the (unmodified) query is executed directly on the precomputed result. This approach, called **view materialization**, is likely to be much faster than the query modification approach because the complex view need not be evaluated when the query is computed. Materialized views can be used during query processing in the same way as regular relations; for example, we can create indexes on materialized views to further speed up query processing. The drawback, of course, is that we must maintain the consistency of the precomputed (or *materialized*) view whenever the underlying tables are updated.

25.9.1 Issues in View Materialization

Three questions must be considered with regard to view materialization:

1. What views should we materialize and what indexes should we build on the materialized views?
2. Given a query on a view and a set of materialized views, can we exploit the materialized views to answer the query?
3. How should we synchronize materialized views with changes to the underlying tables? The choice of synchronization technique depends on several factors, such as whether the underlying tables are in a remote database. We discuss this issue in Section 25.10.

The answers to the first two questions are related. The choice of views to materialize and index is governed by the expected workload, and the discussion of indexing in Chapter 20 is relevant to this question as well. The choice of views to materialize is more complex than just choosing indexes on a set of database tables, however, because the range of alternative views to materialize is wider. The goal is to materialize a small, carefully chosen set of views that can be utilized to quickly answer most of the important queries. Conversely, once we have chosen a set of views to materialize, we have to consider how they can be used to answer a given query.

Consider the `RegionalSales` view. It involves a `JOIN` of `Sales`, `Products`, and `Locations` and is likely to be expensive to compute. On the other hand, if it

is materialized and stored with a clustered B+ tree index on the search key (category, state, sales), we can answer the example query by an index-only search.

Given the materialized view and this index, we can also answer queries of the following form efficiently:

```
SELECT  R.state, SUM (R.sales)
FROM    RegionalSales R
WHERE   R.category = 'Laptop'
GROUP BY R.state
```

To answer such a query, we can use the index on the materialized view to locate the first index leaf entry with *category* = 'Laptop' and then scan the leaf level until we come to the first entry with *category* not equal to Laptop.

The given index is less effective on the following query, for which we are forced to scan the entire leaf level:

```
SELECT  R.state, SUM (R.sales)
FROM    RegionalSales R
WHERE   R.state = 'Wisconsin'
GROUP BY R.category
```

This example indicates how the choice of views to materialize and the indexes to create are affected by the expected workload. This point is illustrated further by our next example.

Consider the following two queries:

```
SELECT  P.category, SUM (S.sales)
FROM    Products P, Sales S
WHERE   P.prod = S.prod
GROUP BY P.category
```

```
SELECT  L.state, SUM (S.sales)
FROM    Locations L, Sales S
WHERE   L.locid = S.locid
GROUP BY L.state
```

These two queries require us to join the Sales table (which is likely to be very large) with another table and aggregate the result. How can we use materialization to speed up these queries? The straightforward approach is to precompute

each of the joins involved (Products with Sales and Locations with Sales) or to precompute each query in its entirety. An alternative approach is to define the following view:

```
CREATE VIEW rrotalSaJes (pid, locid, total)
AS SELECT S.pid, S.locid, SUM (S.sales)
FROM Sales S
GROUP BY S.pid, S.locid
```

The view TotalSales can be materialized and used instead of Sales in our two example queries:

```
SELECT P.category, SUM (T.total)
FROM Products P, TotalSales T
WHERE P.pid = T.pid
GROUP BY P.category
```

```
SELECT L.state, SUM (T.total)
FROM Locations L, TotalSales T
WHERE L.locid = T.locid
GROUP BY L.state
```

25.10 MAINTAINING MATERIALIZED VIEWS

A materialized view is said to be refreshed when we make it consistent with changes to its underlying tables. The process of refreshing a view to keep it consistent with changes to the underlying table is often referred to as view maintenance. Two questions to consider are

1. *How* do we refresh a view when an underlying table is modified? Two issues of particular interest are how to maintain views *incrementally*, that is, without recomputing from scratch when there is a change to an underlying table; and how to maintain views in a distributed environment such as a data warehouse.
2. *When* should we refresh a view in response to a change to an underlying table?

25.10.1 Incremental View Maintenance

A straightforward approach to refreshing a view is to simply recompute the view when an underlying table is modified. This may, in fact, be a reasonable strategy in some cases. For example, if the underlying tables are in a

remote database, the view can be periodically recomputed and sent to the data warehouse where the view is materialized. This has the advantage that the underlying tables need not be replicated at the warehouse.

Whenever possible, however, algorithms for refreshing a view should be **incremental**, in that the cost is proportional to the extent of the change rather than the cost of recomputing the view from scratch.

To understand the intuition behind incremental view maintenance algorithms, observe that a given row in the materialized view can appear several times, depending on how often it was derived. (Recall that duplicates are not eliminated from the result of an SQL query unless the `DISTINCT` clause is used. In this section, we discuss multiset semantics, even when relational algebra notation is used.) The main idea behind incremental maintenance algorithms is to efficiently compute changes to the rows of the view, either new rows or changes to the count associated with a row; if the count of a row becomes 0, the row is deleted from the view.

We present an incremental maintenance algorithm for views defined using projection, binary join, and aggregation; we cover these operations because they illustrate the main ideas. The approach can be extended to other operations such as selection, union, intersection, and (multiset) difference, as well as expressions containing several operators. The key idea is still to maintain the number of derivations for each view row, but the details of how to efficiently compute the changes in view rows and associated counts differ.

Projection Views

Consider a view V defined in terms of a projection on a table R ; that is, $V = \pi(R)$. Every row v in V has an associated count, corresponding to the number of times it can be derived, which is the number of rows in R that yield v when the projection is applied. Suppose we modify R by inserting a collection of rows R_i and deleting a collection of existing rows R_d .¹ We compute $\pi(R_i)$ and add it to V . If the multiset $\pi(R_i)$ contains a row r with count c and r does not appear in V , we add it to V with count c . If r is in V , we add c to its count. We also compute $\pi(R_d)$ and subtract it from V . (Observe that if r appears in $\pi(R_d)$ with count c , it must also appear in V with a higher count;² we subtract c from r 's count in V .)

¹These collections can be multisets of rows. We can treat a row modification as an insert followed by a delete, for simplicity.

²As a simple exercise, consider why this must be so.

As an example, consider the view $\pi_{sales}(Sales)$ and the instance of Sales shown in Figure 25.2. Each row in the view has a single column; the value 25 appears with count 1, and the value 10 appears with count 3. If we delete one of the rows in Sales with sales 10, the count of the (row with) value 10 in the view becomes 2. If we insert a new row into Sales with sales 99, the view now has a row with value 99.

An important point is that we have to maintain the counts associated with rows even if the view definition uses the DISTINCT clause, meaning that duplicates are eliminated from the view. Consider the same view with set semantics—the DISTINCT clause is used in the SQL view definition—and suppose that we delete one of the rows in Sales with sales 10. Does the view now contain a row with value 10? To determine that the answer is yes, we need to maintain the row counts, even though each row (with a nonzero count) is displayed only once in the materialized view.

Join Views

Next, consider a view V defined as a join of two tables, $R \bowtie S$. Suppose we modify R by inserting a collection of rows R_i and deleting a collection of rows R_d . We compute $R_i \bowtie S$ and add the result to V . We also compute $R_d \bowtie S$ and subtract the result from V . Observe that if r appears in $R_d \bowtie S$ with count c , it must also appear in V with a higher count.³

Views with Aggregation

Consider a view V defined over R using GROUP BY on column G and an aggregate operation on column A . Each row v in the view summarizes a group of tuples in R and is of the form $(g, \text{summary})$, where g is the value of the grouping column G and the summary information depends on the aggregate operation. To maintain such a view incrementally, in general, we have to keep a more detailed summary than just the information included in the view. If the aggregate operation is COUNT, we need to maintain only a count c for each row v in the view. If a row r is inserted into R , and there is no row v in V with $v.G = r.G$, we add a new row $(r.G, 1)$. If there is a row v with $v.G = r.G$, we increment its count. If a row r is deleted from R , we decrement the count for the row v with $v.G = r.G$; v can be deleted if its count becomes 0, because then the last row in this group has been deleted from R .

If the aggregate operation is SUM, we have to maintain a sum s and also a count c . If a row r is inserted into R and there is no row v in V with $v.G = r.G$,

³As another simple exercise, consider why this must be so.

we add a new row $\langle r.G, a, 1 \rangle$. If there is a row $\langle r.G, s, c \rangle$, we replace it by $\langle r.G, s + a, c + 1 \rangle$. If a row r is deleted from R , we replace the row $\langle r.G, s, c \rangle$ with $\langle r.G, s - a, c - 1 \rangle$; v can be deleted if its count becomes 0. Observe that without the count, we do not know when to delete v , since the sum for a group could be 0 even if the group contains some rows.

If the aggregate operation is AVG, we have to maintain a sum s , a count c , and the average for each row in the view. The sum and count are maintained incrementally as already described, and the average is computed as s/c .

The aggregate operations MIN and MAX are potentially expensive to maintain. Consider MIN. For each group in R , we maintain (g, m, c) , where m is the minimum value for column A in the group g , and c is the count of the number of rows r in R with $r.G = g$ and $r.A = m$. If a row r is inserted into R and $r.G = g$, if $r.A$ is greater than the minimum m for group g , we can ignore r . If $r.A$ is equal to the minimum m for r 's group, we replace the summary row for the group with $(g, m, c + 1)$. If $r.A$ is less than the minimum m for r 's group, we replace the summary row for the group with $(g, r.A, 1)$. If a row r is deleted from R and $r.A$ is equal to the minimum m for r 's group, then we must decrement the count for the group. If the count is greater than 0, we simply replace the summary row for the group with $(g, m, c - 1)$. However, if the count becomes 0, this means the last row with the recorded minimum A value has been deleted from R and we have to retrieve the smallest A value among the remaining rows in R with group value $r.G$ and this might require retrieval of all rows in R with group value $r.G$.

25.10.2 Maintaining Warehouse Views

The views materialized in a data warehouse can be based on source tables in remote databases. The asynchronous replication techniques discussed in Section 22.11.2 allow us to communicate changes at the source to the warehouse, but refreshing views incrementally in a distributed setting presents some unique challenges. To illustrate this, we consider a simple view that identifies suppliers of Toys.

```
CREATE VIEW ToySuppliers (sid)
AS SELECT S.sid
FROM Suppliers S, Products P
WHERE S.pid = P.pid AND P.category = 'Toys'
```

Suppliers is a new table introduced for this example; let us assume that it has just two fields, *sid* and *pid*, indicating that supplier *sid* supplies part *pid*. The location of the tables Products and Suppliers and the view ToySuppliers

influences how we maintain the view. Suppose that all three are maintained at a single site. We can maintain the view incrementally using the techniques discussed in Section 25.10.1. If a replica of the view is created at another site, we can monitor changes to the materialized view and apply them at the second site using the asynchronous replication techniques from Section 22.11.2.

But, what if Products and Suppliers are at one site and the view is materialized (only) at a second site? To motivate this scenario, we observe that, if the first site is used for operational data and the second site supports complex analysis, the two sites may well be administered by different groups. The option of materializing ToysSuppliers (a view of interest to the second group) at the first site (run by a different group) is not attractive and may not even be possible; the administrators of the first site may not want to deal with someone else's views, and the administrators of the second site may not want to coordinate with someone else whenever they modify view definitions. As another motivation for materializing views at a different location from source tables, observe that Products and Suppliers may be at two different sites. Even if we materialize ToysSuppliers at one of these sites, one of the two source tables is remote.

Now that we have presented motivation for maintaining ToysSuppliers at a location (say, Warehouse) different from the one (say, Source) that contains Products and Suppliers, let us consider the difficulties posed by data distribution. Suppose that a new Products record (with *category* = 'Toys') is inserted. We could try to maintain the view incrementally as follows:

1. The Warehouse site sends this update to the Source site.
2. To refresh the view, we need to check the Suppliers table to find suppliers of the item, and so the Warehouse site asks the Source site for this information.
3. The Source site returns the set of suppliers for the sold item, and the Warehouse site incrementally refreshes the view.

This works when there are no additional changes at the Source site in between steps (1) and (3). If there are changes, however, the materialized view can become incorrect—reflecting a state that can never arise except for anomalies introduced by the preceding, naive, incremental refresh algorithm. To see this, suppose that Products is empty and Suppliers contains just the row $\langle s1, 5 \rangle$ initially, and consider the following sequence of events:

1. Product $pid = 5$ is inserted with *category* = 'Toys'; Source notifies Warehouse.
2. Warehouse asks Source for suppliers of product $pid = 5$. (*The only such supplier at this instant is 81.*)

3. The row (82,5) is inserted into Suppliers; Source notifies \Varehouse.
4. To decide whether 82 should be added to the view, we need to know the category of product $pid = 5$, and \Varehouse asks Source. (*Warehouse has not received an answer to its previous question.*)
5. Source now processes the first query from Warehouse, finds two suppliers for part 5, and returns this information to Warehouse.
6. Warehouse gets the answer to its first question: suppliers 81 and 82, and adds these to the view, each with count 1.
7. Source processes the second query from \Varehouse and responds with the information that part 5 is a toy.
8. Warehouse gets the answer to its second question and accordingly increments the count for supplier 82 in the view.
9. Product $pid = 5$ is now deleted; Source notifies Warehouse.
10. Since the deleted part is a toy, Warehouse decrements the counts of matching view tuples; 81 has count 0 and is removed, but s2 has count 1 and is retained.

Clearly, 82 should not remain in the view after part 5 is deleted. This example illustrates the added subtleties of incremental view maintenance in a distributed environment, and this is a topic of ongoing research.

25.10.3 When Should We Synchronize Views?

A view maintenance policy is a decision about when a view is refreshed, independent of whether the refresh is incremental or not. A view can be refreshed within the same transaction that updates the underlying tables. This is called immediate view maintenance. The update transaction is slowed by the refresh step, and the impact of refresh increases with the number of materialized views that depend on the updated table.

Alternatively, we can defer refreshing the view. Updates are captured in a log and applied subsequently to the materialized view. There are several deferred view maintenance policies:

1. Lazy: The materialized view V is refreshed at the time a query is evaluated using V , if V is not already consistent with its underlying base tables. This approach slows down queries rather than updates, in contrast to immediate view maintenance.

Views for Decision Support: DBMS vendors are enhancing their main relational products to support decision support queries. IBM DB2 supports materialized views with transaction-consistent or user-invoked maintenance. Microsoft SQL Server supports partition views, which are unions of (many) horizontal partitions of a table. These are aimed at a warehousing environment where each partition could be, for example, a monthly update. Queries on partition views are optimized so that only relevant partitions are accessed. Oracle 9i supports materialized views with transaction-consistent, user-invoked, or time-scheduled maintenance.

2. **Periodic:** The materialized view is refreshed periodically, say, once a day. The discussion of the Capture and Apply steps in asynchronous replication (see Section 22.11.2) should be reviewed at this point, since it is very relevant to periodic view maintenance. In fact, many vendors are extending their asynchronous replication features to support materialized views. Materialized views that are refreshed periodically are also called snapshots.
3. **Forced:** The materialized view is refreshed after a certain number of changes have been made to the underlying tables.

In periodic and forced view maintenance, queries may see an instance of the materialized view that is not consistent with the current state of the underlying tables. That is, the queries would see a different set of rows if the view definition was recomputed. This is the price paid for fast updates and queries, and the trade-off is similar to the trade-off made in using asynchronous replication.

25.11 REVIEW QUESTIONS

Answers to the review questions can be found in the listed sections.

- What are *decision support* applications? Discuss the relationship of *complex SQL queries*, *OLAP*, *data mining*, and *data warehousing*. (Section 25.1)
- Describe the multidimensional data model. Explain the distinction between *measures* and *dimensions* and between *fact tables* and *dimension tables*. What is a *star schema*? (Sections 25.2 and 25.2.1)
- Common OLAP operations have received special names: *roll-up*, *drill-down*, *pivoting*, *slicing*, and *dicing*. Describe each of these operations and illustrate them using examples. (Section 25.3)
- Describe the SQL:1999 ROLLUP and CUBE features and their relationship to the OLAP operations. (Section 25.3.1)

- Describe the SQL:1999 WINDOW feature, in particular, framing and ordering of windows. How does it support queries over ordered data? Give examples of queries that are hard to express without this feature. (Section 25.4)
- New query paradigms include *top N queries* and *online aggregation*. Explain the motivation behind these concepts and illustrate them through examples. (Section 25.5)
- Index structures that are especially suitable for OLAP systems include *bitmap indexes* and *join indexes*. Describe these structures. How are bitmap indexes related to B+ trees? (Section 25.6)
- Information about daily operations of an organization is stored in *operational databases*. Why is a *data warehouse* used to store data from operational databases? What issues arise in data warehousing? Discuss *data extraction*, *cleaning*, *transformation*, and *loading*. Discuss the challenges in efficiently *partitioning* and *purging* data. (Section 25.7)
- Why are views important in decision support environments? How are views related to data warehousing and OLAP? Explain the *query modification* technique for answering queries over views and discuss why this is not adequate in decision support environments. (Section 25.8)
- What are the main issues to consider in maintaining materialized views? Discuss how to select views to materialize and how to use materialized views to answer a query. (Section 25.9)
- How can views be maintained *incrementally*? Discuss all the relational algebra operators and aggregation. (Section 25.10.1)
- Use an example to illustrate the added complications for incremental view maintenance introduced by data distribution. (Section 25.10.2)
- Discuss the choice of an appropriate *maintenance policy* for when to refresh a view. (Section 25.10.3)

EXERCISES

Exercise 25.1 Briefly answer the following questions:

1. How do warehousing, OLAP, and data mining complement each other?
2. What is the relationship between data warehousing and data replication? Which form of replication (synchronous or asynchronous) is better suited for data warehousing? Why?
3. What is the role of the metadata repository in a data warehouse? How does it differ from a catalog in a relational DBMS?
4. What considerations are involved in designing a data warehouse?

5. Once a warehouse is designed and loaded, how is it kept current with respect to changes to the source databases?
6. One of the advantages of a warehouse is that we can use it to track how the contents of a relation change over time; in contrast, we have only the current snapshot of a relation in a regular DBMS. Discuss how you would maintain the history of a relation R , taking into account that 'old' information must somehow be purged to make space for new information.
7. Describe dimensions and measures in the multidimensional data model.
8. What is a fact table, and why is it so important from a performance standpoint?
9. What is the fundamental difference between MOLAP and ROLAP systems?
10. What is a star schema? Is it typically in BCNF? Why or why not?
11. How is data mining different from OLAP?

Exercise 25.2 Consider the instance of the Sales relation shown in Figure 25.2.

1. Show the result of pivoting the relation on *pid* and *timeid*.
2. Write a collection of SQL queries to obtain the same result as in the previous part.
3. Show the result of pivoting the relation on *pid* and *lacid*.

Exercise 25.3 Consider the cross-tabulation of the Sales relation shown in Figure 25.5.

1. Show the result of roll-up on *lacid* (i.e., state).
2. Write a collection of SQL queries to obtain the same result as in the previous part.
3. Show the result of roll-up on *lacid* followed by drill-down on *pid*.
4. Write a collection of SQL queries to obtain the same result as in the previous part, starting with the cross-tabulation shown in Figure 25.5.

Exercise 25.4 Briefly answer the following questions:

1. What are the differences between the WINDOW clause and the GROUP BY clause?
2. Give an example query that cannot be expressed in SQL without the WINDOW clause but that can be expressed with the WINDOW clause.
3. What is the *frame* of a window in SQL:1999?
4. Consider the following simple GROUP BY query.

```
SELECT  T.year, SUM (S.sales)
FROM    Sales S, Times T
WHERE   S.timeid=T.timeid
GROUP BY T.year
```

Can you write this query in SQL:1999 without using a GROUP BY clause? (Hint: Use the SQL:1999 WINDOW clause.)

Exercise 25.5 Consider the Locations, Products, and Sales relations shown in Figure 25.2. Write the following queries in SQL:1999 using the WINDOW clause whenever you need it.

1. Find the percentage change in the total monthly sales for each location.
2. Find the percentage change in the total quarterly sales for each product.

3. Find the average daily sales over the preceding 30 days for each product.
4. For each week, find the maximum moving average of sales over the preceding four weeks.
5. Find the top three locations ranked by total sales.
6. Find the top three locations ranked by cumulative sales, for every month over the past year.
7. Rank all locations by total sales over the past year, and for each location print the difference in total sales relative to the location behind it.

Exercise 25.6 Consider the Customers relation and the bitmap indexes shown in Figure 25.9.

1. For the same data, if the underlying set of rating values is assumed to range from 1 to 10, show how the bitmap indexes would change.
2. How would you use the bitmap indexes to answer the following queries? If the bitmap indexes are not useful, explain why.
 - (a) How many customers with a rating less than 3 are male?
 - (b) What percentage of customers are male?
 - (c) How many customers are there?
 - (d) How many customers are named Woo?
 - (e) Find the rating value with the greatest number of customers and also find the number of customers with that rating value; if several rating values have the maximum number of customers, list the requested information for all of them. (Assume that very few rating values have the same number of customers.)

Exercise 25.7 In addition to the Customers table of Figure 25.9 with bitmap indexes on *gender* and *rating*, assume that you have a table called Prospects, with fields *rating* and *prospectid*. This table is used to identify potential customers.

1. Suppose that you also have a bitmap index on the *rating* field of Prospects. Discuss whether or not the bitmap indexes would help in computing the join of Customers and Prospects on *rating*.
2. Suppose you have *no* bitmap index on the *rating* field of Prospects. Discuss whether or not the bitmap indexes on Customers would help in computing the join of Customers and Prospects on *rating*.
3. Describe the use of a join index to support the join of these two relations with the join condition *custid=prospectid*.

Exercise 25.8 Consider the instances of the Locations, Products, and Sales relations shown in Figure 25.2.

1. Consider the basic join indexes described in Section 25.6.2. Suppose you want to optimize for the following two kinds of queries: Query 1 finds sales in a given city, and Query 2 finds sales in a given state. Show the indexes you would create on the example instances shown in Figure 25.2.
2. Consider the bitmapped join indexes described in Section 25.6.2. Suppose you want to optimize for the following two kinds of queries: Query 1 finds sales in a given city, and Query 2 finds sales in a given state. Show the indexes that you would create on the example instances shown in Figure 25.2.

3. Consider the basic join indexes described in Section 25.6.2. Suppose you want to optimize for these two kinds of queries: Query 1 finds sales in a given city for a given product name, and Query 2 finds sales in a given state for a given product category. Show the indexes that you would create on the example instances shown in Figure 25.2.
4. Consider the bitmapped join indexes described in Section 25.6.2. Suppose you want to optimize for these two kinds of queries: Query 1 finds sales in a given city for a given product name, and Query 2 finds sales in a given state for a given product category. Show the indexes that you would create on the example instances shown in Figure 25.2.

Exercise 25.9 Consider the view NumReservations defined as:

```
CREATE VIEW NumReservations (sid, sname, nres)
AS SELECT S.sid, S.sname, COUNT (*)
   FROM   Sailors S, Reserves R
   WHERE  S.sid = R.sid
   GROUP BY S.sid, S.sname
```

1. How is the following query, which is intended to find the highest number of reservations made by some one sailor, rewritten using query modification?

```
SELECT  MAX (N.numres)
FROM    NumReservations N
```

2. Consider the alternatives of computing on demand and view materialization for the preceding query. Discuss the pros and cons of materialization.
3. Discuss the pros and cons of materialization for the following query:

```
SELECT  N.sname, MAX (N.numres)
FROM    NumReservations N
GROUP BY N.sname
```

Exercise 25.10 Consider the Locations, Products, and Sales relations in Figure 25.2.

1. To decide whether to materialize a view, what factors do we need to consider?
2. Assume that we have defined the following materialized view:

```
SELECT  L.state, S.sales
FROM    Locations L, Sales S
WHERE   S.locid=L.locid
```

- (a) Describe what auxiliary information the algorithm for incremental view maintenance from Section 25.10.1 maintains and how this data helps in maintaining the view incrementally.
- (b) Discuss the pros and cons of materializing this view.
3. Consider the materialized view in the previous question. Assume that the relations Locations and Sales are stored at one site, but the view is materialized on a second site. Why would we ever want to maintain the view at a second site? Give a concrete example where the view could become inconsistent.
4. Assume that we have defined the following materialized view:

```
SELECT  T.year, L.state, SUM (S.sales)
FROM    Sales S, Times T, Locations L
WHERE   S.timeid=T.timeid AND S.locid=L.locid
GROUP BY T.year, L.state
```

- (a) Describe what auxiliary information the algorithm for incremental view maintenance from Section 25.10.1 maintains, and how this data helps in maintaining the view incrementally.
- (b) Discuss the pros and cons of materializing this view.

BIBLIOGRAPHIC NOTES

A good survey of data warehousing and OLAP is presented in [161], which is the source of Figure 25.10. [686] provides an overview of OLAP and statistical database research, showing the strong parallels between concepts and research in these two areas. The book by Kimball [436], one of the pioneers in warehousing, and the collection of papers in [2] offer a good practical introduction to the area. The term OLAP was popularized by Codd's paper [191]. For a recent discussion of the performance of algorithms utilizing bitmap and other nontraditional index structures, see [575].

Stonebraker discusses how queries on views can be converted to queries on the underlying tables through query modification [713]. Hanson compares the performance of query modification versus immediate and deferred view maintenance [365]. Srivastava and Roterl present an analytical model of materialized view maintenance algorithms [707]. A number of papers discuss how materialized views can be incrementally maintained as the underlying relations are changed. Research into this area has become very active recently, in part because of the interest in *data warehouses*, which can be thought of as collections of views over relations from various sources. An excellent overview of the state of the art can be found in [348], which contains a number of influential papers together with additional material that provides context and background. The following partial list should provide pointers for further reading: [100, 192, 193, 349, 369, 570, 601, 635, 664, 705, 800].

Gray et al. introduced the CUBE operator [335], and optimization of CUBE queries and efficient maintenance of the result of a CUBE query have been addressed in several papers, including [12, 94, 216, 367, 380, 451, 634, 638, 687, 799]. Related algorithms for processing queries with aggregates and grouping are presented in [160, 166]. Rao, Badia, and Van Gucht address the implementation of queries involving generalized quantifiers such as *a majority of* [618]. Srivastava, Tan, and Lum describe an access method to support processing of aggregate queries [708]. Shannugasundaran et al. discuss how to maintain compressed cubes for approximate answering of aggregate queries in [675].

SQL:1999's support for OLAP, including CUBE and WINDOW constructs, is described in [523]. The windowing extensions are very similar to SQL extension for querying sequence data, called SRQL, proposed in [610]. Sequence queries have received a lot of attention recently. Extending relational systems, which deal with sets of records, to deal with sequences of records is investigated in [473, 665, 671].

There has been recent interest in one-pass query evaluation algorithms and database management for data streams. A recent survey of data management for data streams and algorithms for data stream processing can be found in [49]. Examples include quantile and order-statistics computation [340, 506], estimating frequency moments and join sizes [34, 35], estimating correlated aggregates [310], multidimensional regression analysis [173], and computing one-dimensional (i.e., single-attribute) histograms and Haar wavelet decompositions [319, 345].

Other work includes techniques for incrementally maintaining equi-depth histograms [313] and Haar wavelets [515], maintaining samples and simple statistics over sliding windows [201],

as well as general, high-level architectures for stream database systems [50]. Zdonik et al. describe the architecture of a database system for monitoring data streams [795]. A language infrastructure for developing data stream applications is described by Cortes et al. [199].

Carey and Kossmann discuss how to evaluate queries for which only the first few answers are desired [135, 136]. Donjerkovic and Ramanakrishnan consider how a probabilistic approach to query optimization can be applied to this problem [229]. [120] compares several strategies for evaluating Top N queries. Hellerstein et al. discuss how to return approximate answers to aggregate queries and to refine them ‘online.’ [47, 374]. This work has been extended to online computation of joins [354], online reordering [617] and to adaptive query processing [48].

There has been recent interest in approximate query answering, where a small synopsis data structure is used to give fast approximate query answers with provable performance guarantees [7, 8, 61, 159, 167, 314, 759].



26

DATA MINING

- ☛ What is data mining?
- ☛ What is market basket analysis? What algorithms are efficient for counting co-occurrences?
- ☛ What is the a priori property and why is it important?
- ☛ What is a Bayesian network?
- ☛ What is a classification rule? What is a regression rule?
- ☛ What is a decision tree? How are decision trees constructed?
- ☛ What is clustering? What is a simple clustering algorithm?
- ☛ What is a similarity search over sequences? How is it implemented?
- ☛ How can data mining models be constructed incrementally?
- ☛ What are the new mining challenges presented by data streams?
- Key concepts: data mining, KDD process; market basket analysis, co-occurrence counting, association rule, generalized association rule; decision tree, classification tree; clustering; sequence similarity search; incremental model maintenance, data streams, block evolution

The secret of success is to know something nobody else knows.

—Aristotle Onassis

Data mining consists of finding interesting trends or patterns in large datasets to guide decisions about future activities. There is a general expectation that

data mining tools should be able to identify these patterns in the data with minimal user input. The patterns identified by such tools can give a data analyst useful and unexpected insight that can be more carefully investigated subsequently, perhaps using other decision support tools. In this chapter, we discuss several widely studied data mining tasks. Commercial tools are available for each of these tasks from major vendors, and the area is rapidly growing in importance as these tools gain acceptance in the user community.

We start in Section 26.1 by giving a short introduction to data mining. In Section 26.2, we discuss the important task of counting co-occurring items. In Section 26.3, we discuss how this task arises in data mining algorithms that discover rules from the data. In Section 26.4, we discuss patterns that represent rules in the form of a tree. In Section 26.5, we introduce a different data mining task, called *clustering*, and describe how to find clusters in large datasets. In Section 26.6, we describe how to perform similarity search over sequences. We discuss the challenges in mining evolving data and data streams in Section 26.7. We conclude with a short overview of other data mining tasks in Section 26.8.

26.1 INTRODUCTION TO DATA MINING

Data mining is related to the subarea of statistics called *exploratory data analysis*, which has similar goals and relies on statistical measures. It is also closely related to the subareas of artificial intelligence called *knowledge discovery* and *machine learning*. The important distinguishing characteristic of data mining is that the volume of data is very large; although ideas from these related areas of study are applicable to data mining problems, *scalability with respect to data size* is an important new criterion. An algorithm is scalable if the running time grows (linearly) in proportion to the dataset size, holding the available system resources (e.g., amount of main memory and CPU processing speed) constant. Old algorithms must be adapted or new algorithms developed to ensure scalability when discovering patterns from data.

Finding useful trends in datasets is a rather loose definition of data mining: In a certain sense, all database queries can be thought of as doing just this. Indeed, we have a continuum of analysis and exploration tools with SQL queries at one end, OLAP queries in the middle, and data mining techniques at the other end. SQL queries are constructed using relational algebra (with some extensions), OLAP provides higher-level querying idioms based on the multidimensional data model, and data mining provides the most abstract analysis operations. We can think of different data mining tasks as complex ‘queries’ specified at a high level, with a few parameters that are user-definable, and for which specialized algorithms are implemented.

SQL/MM: Data Mining SQL/MM: The SQL/MM: Data Mining extension of the SQL:1999 standard supports four kinds of data mining models: *frequent itemsets and association rules*, *clusters of records*, *regression trees*, and *classification trees*. Several new data types are introduced. These data types play several roles. Some represent a particular class of model (e.g., `DM_RegressionModel`, `DM_ClusteringModel`); some specify the input parameters for a mining algorithm (e.g., `DM_RegTask`, `DM_ClusTask`); some describe the input data (e.g., `DM_LogicalDataSpec`, `DM_MiningData`); and some represent the result of executing a mining algorithm (e.g., `DM_RegResult`, `DM_ClusResult`). Taken together, these classes and their methods provide a standard interface to data mining algorithms that can be invoked from any SQL:1999 database system. The data mining models can be exported in a standard XML format called **Predictive Model Markup Language (PMML)**; models represented using PMML can be imported as well.

In the real world, data mining is much more than simply applying one of these algorithms. Data is often noisy or incomplete, and unless this is understood and corrected for, it is likely that many interesting patterns will be missed and the reliability of detected patterns will be low. Further, the analyst must decide what kinds of mining algorithms are called for, apply them to a well-chosen subset of data samples and variables (i.e., tuples and attributes), digest the results, apply other decision support and mining tools, and iterate the process.

26.1.1 The Knowledge Discovery Process

The knowledge discovery and data mining (KDD) process can roughly be separated into four steps.

1. **Data Selection:** The target subset of data and the attributes of interest are identified by examining the entire raw dataset.
2. **Data Cleaning:** Noise and outliers are removed, field values are transformed to common units and some new fields are created by combining existing fields to facilitate analysis. The data is typically put into a relational format, and several tables might be combined in a *denormalization* step.
3. **Data Mining:** We apply data mining algorithms to extract interesting patterns.
4. **Evaluation:** The patterns are presented to end-users in an understandable form, for example, through visualization.

The results of any step in the KDD process might lead us back to an earlier step to redo the process with the new knowledge gained. In this chapter, however, we limit ourselves to looking at algorithms for some specific data mining tasks. We do not discuss other aspects of the KDD process.

26.2 COUNTING CO-OCCURRENCES

We begin by considering the problem of counting co-occurring items, which is motivated by problems such as market basket analysis. A market basket is a collection of items purchased by a customer in a single customer transaction. A customer transaction consists of a single visit to a store, a single order through a mail-order catalog, or an order at a store on the Web. (In this chapter, we often abbreviate *customer transaction* to *transaction* when there is no confusion with the usual meaning of *transaction* in a DBMS context, which is an execution of a user program.) A common goal for retailers is to identify items that are purchased together. This information can be used to improve the layout of goods in a store or the layout of catalog pages.

<i>transid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
111	201	5/1/99	pen	2
111	201	5/1/99	ink	1
111	201	5/1/99	milk	3
111	201	5/1/99	juice	6
112	105	6/3/99	pen	1
112	105	6/3/99	ink	1
112	105	6/3/99	milk	1
113	106	5/10/99	pen	1
113	106	5/10/99	ink	1
114	201	6/1/99	pen	2
114	201	6/1/99	ink	2
114	201	6/1/99	juice	4
114	201	6/1/99	water	1

Figure 26.1 The Purchases Relation

26.2.1 Frequent Itemsets

We use the Purchases relation shown in Figure 26.1 to illustrate frequent itemsets. The records are shown sorted into groups by transaction. All tuples in a group have the same *transid*, and together they describe a customer transaction, which involves purchases of one or more items. A transaction occurs

on a given date, and the name of each purchased item is recorded, along with the purchased quantity. Observe that there is redundancy in Purchases: It can be decomposed by storing *transid-custid-date* triples in a separate table and dropping *custid* and *date* from Purchases; this may be how the data is actually stored. However, it is convenient to consider the Purchases relation, as shown in Figure 26.1, to compute frequent itemsets. Creating such ‘denormalized’ tables for ease of data mining is commonly done in the data cleaning step of the EDD process.

By examining the set of transaction groups in Purchases, we can make observations of the form: “In 75% of the transactions a pen and ink are purchased together.” This statement describes the transactions in the database. Extrapolation to future transactions should be done with caution, as discussed in Section 26.3.6. Let us begin by introducing the terminology of market basket analysis. An itemset is a set of items. The support of an itemset is the fraction of transactions in the database that contain all the items in the itemset. In our example, the itemset {pen, ink} has 75% support in Purchases. We can therefore conclude that pens and ink are frequently purchased together. If we consider the itemset {milk, juice}, its support is only 25%; milk and juice are not purchased together frequently.

Usually the number of sets of items frequently purchased together is relatively small, especially as the size of the itemsets increases. We are interested in all itemsets whose support is higher than a user-specified minimum support called *minsup*; we call such itemsets frequent itemsets. For example, if the minimum support is set to 70%, then the frequent itemsets in our example are {pen}, {ink}, {milk}, {pen, ink}, and {pen, milk}. Note that we are also interested in itemsets that contain only a single item since they identify frequently purchased items.

We show an algorithm for identifying frequent itemsets in Figure 26.2. This algorithm relies on a simple yet fundamental property of frequent itemsets:

The a Priori Property: Every subset of a frequent itemset is also a frequent itemset.

The algorithm proceeds iteratively, first identifying frequent itemsets with just one item. In each subsequent iteration, frequent itemsets identified in the previous iteration are extended with another item to generate larger candidate itemsets. By considering only itemsets obtained by enlarging frequent itemsets, we greatly reduce the number of candidate frequent itemsets; this optimization is crucial for efficient execution. The a priori property guarantees that this optimization is correct; that is, we do not miss any frequent itemsets. A single scan of all transactions (the Purchases relation in our example) suffices to


```

foreach itemset,                                     Level 1
    check if it is a frequent itemset // appears in > minsup transactions
k = 1
repeat // Iterative, level-wise identification of frequent itemsets
    foreach new frequent itemset Ik with k items // Level k + 1
        generate all itemsets Ik+1 with k + 1 items, Ik ⊆ Ik+1
        Scan all transactions once and check if
        the generated k + 1-itemsets are frequent
        k = k + 1
until no new frequent itemsets are identified

```

Figure 26.2 An Algorithm for Finding Frequent Itemsets

determine which candidate itemsets generated in an iteration are frequent. The algorithm terminates when no new frequent itemsets are identified in an iteration.

We illustrate the algorithm on the Purchases relation in Figure 26.1, with *minsup* set to 70%. In the first iteration (Level 1), we scan the Purchases relation and determine that each of these one-item sets is a frequent itemset: {*pen*} (appears in all four transactions), {*ink*} (appears in three out of four transactions), and {*rnilk*} (appears in three out of four transactions).

In the second iteration (Level 2), we extend each frequent itemset with an additional item and generate the following candidate itemsets: {*pen*, *ink*}, {*pen*, *milk*}, {*pen*, *juice*}, {*ink*, *rnilk*}, {*ink*, *juice*}, and {*rnilk*, *juice*}. By scanning the Purchases relation again, we determine that the following are frequent itemsets: {*pen*, *ink*} (appears in three out of four transactions), and {*pen*, *rnilk*} (appears in three out of four transactions).

In the third iteration (Level 3), we extend these itemsets with an additional item and generate the following candidate itemsets: {*pen*, *ink*, *milk*}, {*pen*, *ink*, *juice*}, and {*pen*, *milk*, *juice*}. (Observe that {*ink*, *milk*, *juice*} is not generated.) A third scan of the Purchases relation allows us to determine that none of these is a frequent itemset.

The simple algorithm presented here for finding frequent itemsets illustrates the principal feature of more sophisticated algorithms, namely, the iterative generation and testing of candidate itemsets. We consider one important refinement of this simple algorithm. Generating candidate itemsets by adding an item to a known frequent itemset is an attempt to limit the number of candidate itemsets using the a priori property. The a priori property implies that a can-

candidate itemset can be frequent only if all its subsets are frequent. Thus, we can reduce the number of candidate itemsets further—*a priori*, or before scanning the Purchases database—by checking whether all subsets of a newly generated candidate itemset are frequent. Only if all subsets of a candidate itemset are frequent do we compute its support in the subsequent database scan. Compared to the simple algorithm, this refined algorithm generates fewer candidate itemsets at each level and thus reduces the amount of computation performed during the database scan of Purchases.

Consider the refined algorithm on the Purchases table in Figure 26.1 with $\text{minsup} = 70\%$. In the first iteration (Level 1), we determine the frequent itemsets of size one: $\{pen\}$, $\{ink\}$, and $\{milk\}$. In the second iteration (Level 2), only the following candidate itemsets remain when scanning the Purchases table: $\{pen, ink\}$, $\{pen, milk\}$, and $\{ink, milk\}$. Since $\{juice\}$ is not frequent, the itemsets $\{pen, juice\}$, $\{ink, juice\}$, and $\{milk, juice\}$ cannot be frequent as well and we can eliminate those itemsets *a priori*, that is, without considering them during the subsequent scan of the Purchases relation. In the third iteration (Level 3), no further candidate itemsets are generated. The itemset $\{pen, ink, milk\}$ cannot be frequent since its subset $\{ink, milk\}$ is not frequent. Thus, the improved version of the algorithm does not need a third scan of Purchases.

26.2.2 Iceberg Queries

We introduce iceberg queries through an example. Consider again the Purchases relation shown in Figure 26.1. Assume that we want to find pairs of customers and items such that the customer has purchased the item more than five times. We can express this query in SQL as follows:

```
SELECT  P.custid, P.item, SUM (P.qty)
FROM    Purchases P
GROUP BY P.custid, P.item
HAVING  SUM (P.qty) > 5
```

Think about how this query would be evaluated by a relational DBMS. Conceptually, for each $(custid, item)$ pair, we need to check whether the sum of the qty field is greater than 5. One approach is to make a scan over the Purchases relation and maintain running sums for each $(custid, item)$ pair. This is a feasible execution strategy as long as the number of pairs is small enough to fit into main memory. If the number of pairs is larger than main memory, more expensive query evaluation plans, which involve either sorting or hashing, have to be used.

The query has an important property not exploited by the preceding execution strategy: Even though the Purchases relation is potentially very large and the

number of $(custid, item)$ groups can be huge, the output of the query is likely to be relatively small because of the condition in the HAVING clause. Only groups where the customer has purchased the item more than five times appear in the output. For example, there are nine groups in the query over the Purchases relation shown in Figure 26.1, although the output contains only three records. The number of groups is very large, but the answer to the query—the tip of the iceberg—is usually very small. Therefore, we call such a query an iceberg query. In general, given a relational schema R with attributes A_1, A_2, \dots, A_k , and B and an aggregation function $aggr$, an iceberg query has the following structure:

```
SELECT  R.A1, R.A2, ..., R.Ak, aggr(R.B)
FROM    Relation R,
GROUP BY R.A1, ..., R.Ak
HAVING  aggr(R.B) >= constant
```

Traditional query plans for this query that use sorting or hashing first compute the value of the aggregation function for all groups and then eliminate groups that do not satisfy the condition in the HAVING clause.

Comparing the query with the problem of finding frequent itemsets discussed in the previous section, there is a striking similarity. Consider again the Purchases relation shown in Figure 26.1 and the iceberg query from the beginning of this section. We are interested in $(custid, item)$ pairs that have $SUM(P.qty) > 5$. Using a variation of the a priori property, we can argue that we only have to consider values of the *custid* field where the customer has purchased at least five items. We can generate such items through the following query:

```
SELECT  P.custid
FROM    Purchases P
GROUP BY P.custid
HAVING  SUM(P.qty) > 5
```

Similarly, we can restrict the candidate values for the item field through the following query:

```
SELECT  P.item
FROM    Purchases P
GROUP BY P.item
HAVING  SUM(P.qty) > 5
```

If we restrict the computation of the original iceberg query to $(custid, item)$ groups where the field values are in the output of the previous two queries, we eliminate a large number of $(custid, item)$ pairs a priori. So, a possible

evaluation strategy is to first compute candidate values for the *custid* and *item* fields, and use combinations of only these values in the evaluation of the original iceberg query. We first generate candidate field values for individual fields and use only those values that survive the a priori pruning step as expressed in the two previous queries. Thus, the iceberg query is amenable to the same bottom-up evaluation strategy used to find frequent itemsets. In particular, we can use the a priori property as follows: We keep a counter for a group only if each individual component of the group satisfies the condition expressed in the HAVING clause. The performance improvements of this alternative evaluation strategy over traditional query plans can be very significant in practice.

Even though the bottom-up query processing strategy eliminates many groups a priori, the number of (*custid*, *item*) pairs can still be very large in practice; even larger than main memory. Efficient strategies that use sampling and more sophisticated hashing techniques have been developed; the bibliographic notes at the end of the chapter provide pointers to the relevant literature.

26.3 MINING FOR RULES

Many algorithms have been proposed for discovering various forms of rules that succinctly describe the data. We now look at some widely discussed forms of rules and algorithms for discovering them.

26.3.1 Association Rules

We use the Purchases relation shown in Figure 26.1 to illustrate association rules. By examining the set of transactions in Purchases, we can identify rules of the form:

$$\{pen\} \Rightarrow \{ink\}$$

This rule should be read as follows: “If a pen is purchased in a transaction, it is likely that ink is also be purchased in that transaction.” It is a statement that describes the transactions in the database; extrapolation to future transactions should be done with caution, as discussed in Section 26.3.6. More generally, an association rule has the form $LHS \Rightarrow RHS$, where both *LHS* and *RHS* are sets of items. The interpretation of such a rule is that if every item in *LHS* is purchased in a transaction, then it is likely that the items in *RHS* are purchased as well.

There are two important measures for an association rule:

- **Support:** The support for a set of items is the percentage of transactions that contain all these items. The support for a rule $LHS \Rightarrow RHS$ is the

support for the set of items $LHS \Rightarrow RHS$. For example, consider the rule $\{pen\} \Rightarrow \{ink\}$. The support of this rule is the support of the itemset $\{pen, ink\}$, which is 75%.

- **Confidence:** Consider transactions that contain all items in LHS . The confidence for a rule $LHS \Rightarrow RHS$ is the percentage of such transactions that also contain all items in RHS . More precisely, let $sup(LHS)$ be the percentage of transactions that contain LHS and let $sup(LHS \cup RHS)$ be the percentage of transactions that contain both LHS and RHS . Then the confidence of the rule $LHS \Rightarrow RHS$ is $sup(LHS \cup RHS) / sup(LHS)$. The confidence of a rule is an indication of the strength of the rule. As an example, consider again the rule $\{pen\} \Rightarrow \{ink\}$. The confidence of this rule is 75%; 75% of the transactions that contain the itemset $\{pen\}$ also contain the itemset $\{ink\}$.

26.3.2 An Algorithm for Finding Association Rules

A user can ask for all association rules that have a specified minimum support ($minsup$) and minimum confidence ($minconf$), and various algorithms have been developed for finding such rules efficiently. These algorithms proceed in two steps. In the first step, all frequent itemsets with the user-specified minimum support are computed. In the second step, rules are generated using the frequent itemsets as input. We discussed an algorithm for finding frequent itemsets in Section 26.2; we concentrate here on the rule generation part.

Once frequent itemsets are identified, the generation of all possible candidate rules with the user-specified minimum support is straightforward. Consider a frequent itemset X with support s_x identified in the first step of the algorithm. To generate a rule from X , we divide X into two itemsets, LHS and RHS . The confidence of the rule $LHS \Rightarrow RHS$ is $s_x / sup(LHS)$, the ratio of the support of X and the support of LHS . From the a priori property, we know that the support of LHS is larger than $minsup$, and thus we have computed the support of LHS during the first step of the algorithm. We can compute the confidence values for the candidate rule by calculating the ratio $support(X)/support(LHS)$ and then check how the ratio compares to $minconf$.

In general, the expensive step of the algorithm is the computation of the frequent itemsets, and many different algorithms have been developed to perform this step efficiently. Rule generation—given that all frequent itemsets have been identified—is straightforward.

In the rest of this section, we discuss some generalizations of the problem.

26.3.3 Association Rules and ISA Hierarchies

In many cases, an **ISA** hierarchy or category hierarchy is imposed on the set of items. In the presence of a hierarchy, a transaction contains, for each of its items, implicitly all the item's ancestors in the hierarchy. For example, consider the category hierarchy shown in Figure 26.3. Given this hierarchy, the Purchases relation is conceptually enlarged by the eight records shown in Figure 26.4. That is, the Purchases relation has all tuples shown in Figure 26.1 in addition to the tuples shown in Figure 26.4.

The hierarchy allows us to detect relationships between items at different levels of the hierarchy. As an example, the support of the itemset $\{ink, juice\}$ is 50%, but if we replace *juice* with the more general category *beverage*, the support of the resulting itemset $\{ink, beverage\}$ increases to 75%. In general, the support of an itemset can increase only if an item is replaced by one of its ancestors in the ISA hierarchy.

Assuming that we actually physically add the eight records shown in Figure 26.4 to the Purchases relation, we can use any algorithm for computing frequent itemsets on the augmented database. Assuming that the hierarchy fits into main memory, we can also perform the addition on-the-fly while we scan the database, as an optimization.



Figure 26.3 An ISA Category Taxonomy

<i>transid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
111	201	5/1/99	stationery	3
111	201	5/1/99	beverage	9
112	105	6/3/99	stationery	2
112	105	6/3/99	beverage	1
113	106	5/10/99	stationery	1
113	106	5/10/99	beverage	1
114	201	6/1/99	stationery	4
114	201	6/1/99	beverage	5

Figure 26.4 Conceptual Additions to the Purchases Relation with ISA Hierarchy

26.3.4 Generalized Association Rules

Although association rules have been most widely studied in the context of market basket analysis, or analysis of customer transactions, the concept is more general. Consider the Purchases relation as shown in Figure 26.5, grouped by *custid*. By examining the set of customer groups, we can identify association rules such as $\{\text{pen}\} \Rightarrow \{\text{milk}\}$. This rule should now be read as follows: "If a pen is purchased by a customer, it is likely that milk is also be purchased by that customer." In the Purchases relation shown in Figure 26.5, this rule has both support and confidence of 100%.

<i>transid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
112	105	6/3/99	pen	1
112	105	6/3/99	ink	1
112	105	6/3/99	milk	1
113	106	5/10/99	pen	1
113	106	5/10/99	milk	1
114	201	5/15/99	pen	2
114	201	5/15/99	ink	2
114	201	5/15/99	juice	4
114	201	6/1/99	water	1
111	201	5/1/99	pen	2
111	201	5/1/99	ink	1
111	201	5/1/99	milk	3
111	201	5/1/99	juice	6

Figure 26.5 The Purchases Relation Sorted on Customer ID

Similarly, we can group tuples by date and identify association rules that describe purchase behavior on the same day. As an example consider again the Purchases relation. In this case, the rule $\{\text{pen}\} \Rightarrow \{\text{milk}\}$ is now interpreted as follows: "On a day when a pen is purchased, it is likely that milk is also be purchased."

If we use the *date* field as grouping attribute, we can consider a more general problem called calendric market basket analysis. In calendric market basket analysis, the user specifies a collection of calendars. A calendar is any group of dates, such as *every Sunday in the year 1999*, or *every first of the month*. A rule holds if it holds on every day in the calendar. Given a calendar, we can compute association rules over the set of tuples whose *date* field falls within the calendar.

By specifying interesting calendars, we can identify rules that might not have enough support and confidence with respect to the entire database but have enough support and confidence on the subset of tuples that fall within the calendar. On the other hand, even though a rule might have enough support and confidence with respect to the complete database, it might gain its support only from tuples that fall within a calendar. In this case, the support of the rule over the tuples within the calendar is significantly higher than its support with respect to the entire database.

As an example, consider the Purchases relation with the calendar *every first of the month*. Within this calendar, the association rule $pen \Rightarrow juice$ has support and confidence of 100%, whereas over the entire Purchases relation, this rule only has 50% support. On the other hand, within the calendar, the rule $pen \Rightarrow milk$ has support and confidence of 50%, whereas over the entire Purchases relation it has support and confidence of 75%.

More general specifications of the conditions that must be true within a group for a rule to hold (for that group) have also been proposed. We might want to say that all items in the *LHS* have to be purchased in a quantity of less than two items, and all items in the *RHS* must be purchased in a quantity of more than three.

Using different choices for the grouping attribute and sophisticated conditions as in the preceding examples, we can identify rules more complex than the basic association rules discussed earlier. These more complex rules, nonetheless, retain the essential structure of an association rule as a condition over a group of tuples, with support and confidence measures defined as usual.

26.3.5 Sequential Patterns

Consider the Purchases relation shown in Figure 26.1. Each group of tuples, having the same *custid* value, can be thought of as a *sequence* of transactions ordered by *date*. This allows us to identify frequently arising buying patterns over time.

We begin by introducing the concept of a sequence of itemsets. Each transaction is represented by a set of tuples, and by looking at the values in the *item* column, we get a set of items purchased in that transaction. Therefore, the sequence of transactions associated with a customer corresponds naturally to a sequence of itemsets purchased by the customer. For example, the sequence of purchases for customer 201 is $\langle \{pen, ink, milk, juice\}, \{pen, ink, juice\} \rangle$.

A subsequence of a sequence of itemsets is obtained by deleting one or more itemsets, and is also a sequence of itemsets. We say that a sequence $\langle a_1, \dots, a_m \rangle$ is contained in another sequence S if S has a subsequence $\langle b_1, \dots, b_n \rangle$ such that $a_i \subseteq b_i$, for $1 \leq i \leq m$. Thus, the sequence $\langle \{pen\}, \{ink, milk\}, \{pen, juice\} \rangle$ is contained in $\langle \{pen, link\}, \{shirt\}, \{juice, ink, milk\}, \{juice, pen, milk\} \rangle$. Note that the order of itemsets within each itemset does not matter. However, the order of itemsets does matter: the sequence $\langle \{pen\}, \{ink, milk\}, \{pen, juice\} \rangle$ is not contained in $\langle \{pen, ink\}, \{shirt\}, \{juice, pen, milk\}, \{juice, milk, ink\} \rangle$.

The support for a sequence S of itemsets is the percentage of customer sequences of which S is a subsequence. The problem of identifying sequential patterns is to find all sequences that have a user-specified minimum support. A sequence $\langle a_1, a_2, a_3, \dots, a_m \rangle$ with minimum support tells us that customers often purchase the items in set a_1 in a transaction, then in some subsequent transaction buy the items in set a_2 , then the items in set a_3 in a later transaction, and so on.

Like association rules, sequential patterns are statements about groups of tuples in the current database. Computationally, algorithms for finding frequently occurring sequential patterns resemble algorithms for finding frequent itemsets. Longer and longer sequences with the required minimum support are identified iteratively in a manner very similar to the iterative identification of frequent itemsets.

26.3.6 The Use of Association Rules for Prediction

Association rules are widely used for prediction, but it is important to recognize that such predictive use is not justified without additional analysis or domain knowledge. Association rules describe existing data accurately but can be misleading when used naively for prediction. For example, consider the rule

$$\{pen\} \Rightarrow \{ink\}$$

The confidence associated with this rule is the conditional probability of an ink purchase given a pen purchase *over the given database*; that is, it is a *descriptive* measure. We might use this rule to guide future sales promotions. For example, we might offer a discount on pens to increase the sales of pens and, therefore, also increase sales of ink.

However, such a promotion assumes that pen purchases are good indicators of ink purchases in *future* customer transactions (in addition to transactions in the current database). This assumption is justified if there is a *causal link* between pen purchases and ink purchases; that is, if buying pens causes the buyer to also buy ink. However, we can infer association rules with high support

and confidence in some situations where there is no causal link between *LHS* and *RHS*. For example, suppose that pens are always purchased together with pencils, perhaps because of customers' tendency to order writing instruments together. We would then infer the rule

$$\{pencil\} \Rightarrow \{ink\}$$

with the same support and confidence as the rule

$$\{pen\} \Rightarrow \{ink\}$$

However, there is no causal link between pencils and ink. If we promote pencils, a customer who purchases several pencils due to the promotion has no reason to buy more ink. Therefore, a sales promotion that discounted pencils in order to increase the sales of ink would fail.

In practice, one would expect that, by examining a large database of past transactions (collected over a long time and a variety of circumstances) and restricting attention to rules that occur often (i.e., that have high support), we minimize inferring misleading rules. However, we should bear in mind that misleading, noncausal rules might still be generated. Therefore, we should treat the generated rules as possibly, rather than conclusively, identifying causal relationships. Although association rules do not indicate causal relationships between the *LHS* and *RHS*, we emphasize that they provide a useful starting point for identifying such relationships, using either further analysis or a domain expert's judgment; this is the reason for their popularity.

26.3.7 Bayesian Networks

Finding causal relationships is a challenging task, as we saw in Section 26.3.6. In general, if certain events are highly correlated, there are many possible explanations. For example, suppose that pens, pencils, and ink are purchased together frequently. It might be that the purchase of one of these items (e.g., ink) depends causally on the purchase of another item (e.g., pen). Or it might be that the purchase of one of these items (e.g., pen) is strongly correlated with the purchase of another (e.g., pencil) because of some underlying phenomenon (e.g., users' tendency to think about writing instruments together) that causally influences both purchases. How can we identify the true causal relationships that hold between these events in the real world?

One approach is to consider each possible combination of causal relationships among the variables or events of interest to us and evaluate the likelihood of each combination on the basis of the data available to us. If we think of each combination of causal relationships as a *model* of the real world underlying the

collected data, we can assign a score to each model by considering how consistent it is (in terms of probabilities, with some simplifying assumptions) with the observed data. Bayesian networks are graphs that can be used to describe a class of such models, with one node per variable or event, and arcs between nodes to indicate causality. For example, a good model for our running example of pens, pencils, and ink is shown in Figure 26.6. In general, the number of possible models is exponential in the number of variables, and considering all models is expensive, so some subset of all possible models is evaluated.

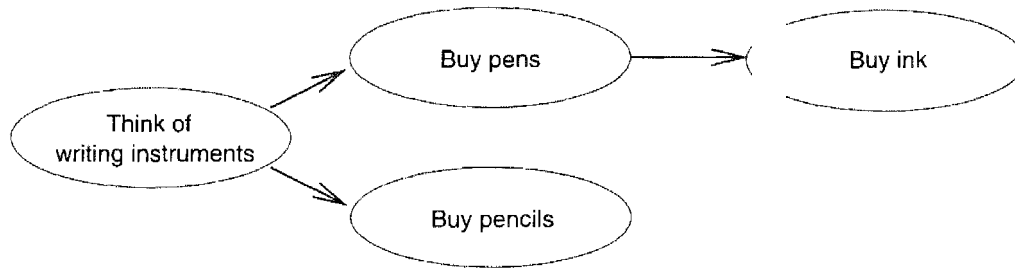


Figure 26.6 Bayesian Network Showing Causality

26.3.8 Classification and Regression Rules

Consider the following view that contains information from a mailing campaign performed by an insurance company:

`InsuranceInfo(age: integer, cartype: string, highrisk: boolean)`

The `InsuranceInfo` view has information about current customers. Each record contains a customer's age and type of car as well as a flag indicating whether the person is considered a high-risk customer. If the flag is true, the customer is considered high-risk. We would like to use this information to identify rules that predict the insurance risk of new insurance applicants whose age and car type are known. For example, one such rule could be: "If *age* is between 16 and 25 and *cartye* is either Sports or Truck, then the risk is high."

Note that the rules we want to find have a specific structure. We are not interested in rules that predict the age or type of car of a person: we are interested only in rules that predict the insurance risk. Thus, there is one designated attribute whose value we wish to predict, and we call this attribute the dependent attribute. The other attributes are called predictor attributes. In our example, the dependent attribute in the `InsuranceInfo` view is the *highrisk* attribute and the predictor attributes are *age* and *cartye*. The general form of the types of rules we want to discover is

$$P_1(X_1) \wedge P_2(X_2) \dots \wedge P_k(X_k) \Rightarrow Y = c$$

The predictor attributes X_1, \dots, X_k are used to predict the value of the dependent attribute Y . Both sides of a rule can be interpreted as conditions on fields of a tuple. The $P_i(X_i)$ are predicates that involve attribute X_i . The form of the predicate depends on the type of the predictor attribute. We distinguish two types of attributes: numerical and categorical. For numerical attributes, we can perform numerical computations, such as computing the average of two values; whereas for categorical attributes, the only allowed operation is testing whether two values are equal. In the InsuranceInfo view, *age* is a numerical attribute whereas *car type* and *highrisk* are categorical attributes. Returning to the form of the predicates, if X_i is a numerical attribute, its predicate P_i is of the form $li \leq X_i \leq hi$; if X_i is a categorical attribute, P_i is of the form $X_i \in \{V_1, \dots, V_j\}$.

If the dependent attribute is categorical, we call such rules classification rules. If the dependent attribute is numerical, we call such rules regression rules.

For example, consider again our example rule: "If *age* is between 16 and 25 and *car type* is either Sports or Truck, then *highrisk* is true." Since *highrisk* is a categorical attribute, this rule is a classification rule. We can express this rule formally as follows:

$$(16 \leq age \leq 25) \wedge (car\ type \in \{Sports, Truck\}) \Rightarrow highrisk = true$$

We can define support and confidence for classification and regression rules, as for association rules:

- **Support:** The support for a condition C is the percentage of tuples that satisfy C . The support for a rule $C1 \Rightarrow C2$ is the support for the condition $C1 \wedge C2$.
- **Confidence:** Consider those tuples that satisfy condition $C1$. The confidence for a rule $C1 \Rightarrow C2$ is the percentage of such tuples that also satisfy condition $C2$.

As a further generalization, consider the right-hand side of a classification or regression rule: $Y = c$. Each rule predicts a value of Y for a given tuple based on the values of predictor attributes X_1, \dots, X_k . We can consider rules of the form

$$P_1(X_1) \wedge \dots \wedge P_k(X_k) \Rightarrow Y = f(X_1, \dots, X_k)$$

where f is some function. We do not discuss such rules further.

Classification and regression rules differ from association rules by considering continuous and categorical fields, rather than only one field that is set-valued. Identifying such rules efficiently presents a new set of challenges; we do not

discuss the general case of discovering such rules. We discuss a special type of such rules in Section 26.4.

Classification and regression rules have many applications. Examples include classification of results of scientific experiments, where the type of object to be recognized depends on the measurements taken; direct mail prospecting, where the response of a given customer to a promotion is a function of his or her income level and age; and car insurance risk assessment, where a customer could be classified as risky depending on age, profession, and car type. Example applications of regression rules include financial forecasting, where the price of coffee futures could be some function of the rainfall in Colombia a month ago, and medical prognosis, where the likelihood of a tumor being cancerous is a function of measured attributes of the tumor.

26.4 TREE-STRUCTURED RULES

In this section, we discuss the problem of discovering classification and regression rules from a relation, but we consider only rules that have a very special structure. The type of rules we discuss can be represented by a tree, and typically the tree itself is the output of the data mining activity. Trees that represent classification rules are called classification trees or decision trees and trees that represent regression rules are called regression trees

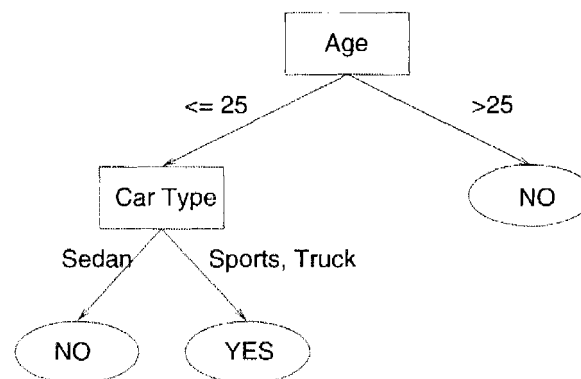


Figure 26.7 Insurance Risk Example Decision Tree

As an example, consider the decision tree shown in Figure 26.7. Each path from the root node to a leaf node represents one classification rule. For example, the path from the root to the leftmost leaf node represents the classification rule: “If a person is 25 years or younger and drives a sedan, then he or she is likely to have a low insurance risk.” The path from the root to the rightmost leaf node represents the classification rule: “If a person is older than 25 years, then he or she is likely to have a low insurance risk.”

Tree-structured rules are very popular since they are easy to interpret. Ease of understanding is very important because the result of any data mining activity needs to be comprehensible by nonspecialists. In addition, studies have shown that, despite limitations in structure, tree-structured rules are very accurate. There exist efficient algorithms to construct tree-structured rules from large databases. We discuss a sample algorithm for decision tree construction in the remainder of this section.

26.4.1 Decision Trees

A decision tree is a graphical representation of a collection of classification rules. Given a data record, the tree directs the record from the root to a leaf. Each internal node of the tree is labeled with a predictor attribute. This attribute is often called a *splitting attribute*, because the data is 'split' based on conditions over this attribute. The outgoing edges of an internal node are labeled with predicates that involve the splitting attribute of the node; every data record entering the node must satisfy the predicate labeling exactly one outgoing edge. The combined information about the splitting attribute and the predicates on the outgoing edges is called the *splitting criterion* of the node. A node with no outgoing edges is called a *leaf node*. Each leaf node of the tree is labeled with a value of the dependent attribute. We consider only binary trees where internal nodes have two outgoing edges, although trees of higher degree are possible.

Consider the decision tree shown in Figure 26.7. The splitting attribute of the root node is *age*, the splitting attribute of the left child of the root node is *car type*. The predicate on the left outgoing edge of the root node is $age \leq 25$, the predicate on the right outgoing edge is $age > 25$.

We can now associate a classification rule with each leaf node in the tree as follows. Consider the path from the root of the tree to the leaf node. Each edge on that path is labeled with a predicate. The conjunction of all these predicates makes up the left-hand side of the rule. The value of the dependent attribute at the leaf node makes up the right-hand side of the rule. Thus, the decision tree represents a collection of classification rules, one for each leaf node.

A decision tree is usually constructed in two phases. In phase one, the *growth phase*, an overly large tree is constructed. This tree represents the records in the input database very accurately; for example, the tree might contain leaf nodes for individual records from the input database. In phase two, the *pruning phase*, the final size of the tree is determined. The rules represented by the tree constructed in phase one are usually overspecialized. By reducing the size of the tree, we generate a smaller number of more general rules that

are better than a very large number of very specialized rules. Algorithms for tree pruning are beyond our scope of discussion here.

Classification tree algorithms build the tree greedily top-down in the following way. At the root node, the database is examined and the locally 'best' splitting criterion is computed. The database is then partitioned, according to the root node's splitting criterion, into two parts, one partition for the left child and one partition for the right child. The algorithm then recurses on each child. This schema is depicted in Figure 26.8.

Input: Node n , partition D , split selection method S

Output: decision tree for D rooted at node n

Top-Down Decision Tree Induction Schema:

BuildTree(Node n , data partition D , split selection method S)

- (1) Apply S to D to find the splitting criterion
- (2) if (a good splitting criterion is found)
- (3) Create two children nodes n_1 and n_2 of n
- (4) Partition D into D_1 and D_2
- (5) BuildTree(n_1 , D_1 , S)
- (6) BuildTree(n_2 , D_2 , S)
- (7) endif

Figure 26.8 Decision Tree Induction Schema

The splitting criterion at a node is found through application of a split selection method. A split selection method is an algorithm that takes as input (part of) a relation and outputs the locally 'best' splitting criterion. In our example, the split selection method examines the attributes *car type* and *age*, selects one of them as splitting attribute, and then selects the splitting predicates. Many different, very sophisticated split selection methods have been developed; the references provide pointers to the relevant literature.

26.4.2 An Algorithm to Build Decision Trees

If the input database fits into main memory, we can directly follow the classification tree induction schema shown in Figure 26.8. How can we construct decision trees when the input relation is larger than main memory? In this case, step (1) in Figure 26.8 fails, since the input database does not fit in memory. But we can make one important observation about split selection methods that helps us to reduce the main memory requirements.

Consider a node of the decision tree. The split selection method has to make two decisions after examining the partition at that node: It has to select the splitting attribute, and it has to select the splitting predicates for the outgo-

<i>age</i>	<i>cartype</i>	<i>highrisk</i>
23	Sedan	false
30	Sports	false
36	Sedan	false
25	Truck	true
30	Sedan	false
23	Truck	true
30	Truck	false
25	Sports	true
18	Sedan	false

Figure 26.9 The InsuranceInfo Relation

ing edges. After selecting the splitting criterion at a node, the algorithm is recursively applied to each of the children of the node. Does a split selection method actually need the complete database partition as input? Fortunately, the answer is no.

Split selection methods that compute splitting criteria that involve a single predictor attribute at each node evaluate each predictor attribute individually. Since each attribute is examined separately, we can provide the split selection method with aggregated information about the database instead of loading the complete database into main memory. Chosen correctly, this aggregated information enables us to compute the same splitting criterion as we would obtain by examining the complete database.

Since the split selection method examines all predictor attributes, we need aggregated information about each predictor attribute. We call this aggregated information the **AVe** set of the predictor attribute. The AVe set of a predictor attribute X at node n is the projection of n 's database partition onto X and the dependent attribute where counts of the individual values in the domain of the dependent attribute are aggregated. (AVC stands for Attribute-Value, Class label, because the values of the dependent attribute are often called class labels.) For example, consider the InsuranceInfo relation as shown in Figure 26.9. The AVe set of the root node of the tree for predictor attribute *age* is the result of the following database query:

```
SELECT  R.age, II.highrisk, COUNT (*)
FROM    InsuranceInfo R
GROUP BY R.age, R.highrisk
```

The AVe set for the left child of the root node for predictor attribute *cartype* is the result of the following query:


```

SELECT  R.carttype, R.highrisk, COUNT (*)
FROM    InsuranceInfo R
WHERE   R.age <= 25
GROUP BY R.carttype, R.highrisk

```

The tVO AVC sets of the root node of the tree are shown in Figure 26.10.

Car type	highrisk	
	true	false
Sedan	0	4
Sports	1	1
Truck	2	1

Age	highrisk	
	true	false
18	0	1
23	1	1
25	2	0
30	0	3
36	0	1

Figure 26.10 AVe Group of the Root Node for the InsuranceInfo Relation

We define the **AVe** group of a node n to be the set of the AVe sets of all predictor attributes at node n . Our example of the InsuranceInfo relation has two predictor attributes; therefore, the AVe group of any node consists of two AVe sets.

How large are AVe sets? Note that the size of the AVe set of a predictor attribute X at node n depends only on the number of distinct attribute values of X and the size of the domain of the dependent attribute. For example, consider the AVe sets shown in Figure 26.10. The AVe set for the predictor attribute *cartype* has three entries, and the AVe set for predictor attribute *age* has five entries, although the InsuranceInfo relation as shown in Figure 26.9 has nine records. For large databases, the size of the AVe sets is independent of the number of tuples in the database, except if there are attributes with very large domains, for example, a real-valued field recorded at a very high precision with many digits after the decimal point.

If we make the simplifying assumption that all the AVe sets of the root node together fit into main memory, then we can construct decision trees for very large databases as follows: We make a scan over the database and construct the AVe group of the root node in memory. Then we run the split selection method of our choice with the AVC group as input. After the split selection method computes the splitting attribute and the splitting predicates on the outgoing nodes, we partition the database and recurse. Note that this algorithm is very similar to the original algorithm shown in Figure 26.8; the only modification necessary is shown in Figure 26.11. In addition, this algorithm is still independent of the actual split selection method involved.

Input: node n , partition D , split selection method S

Output: decision tree for D rooted at node n

Top-Down Decision Tree Induction Schema:

BuHdTree(Node n , data partition D , split selection method S)

(1a) Make a scan over D and construct the AVE group of n in D

(1b) Apply S to the AVE group to find the splitting criterion

Figure 26.11 Classification Tree Induction Refinement with AVE Groups

26.5 CLUSTERING

In this section we discuss the clustering problem. The goal is to partition a set of records into groups such that records within a group are similar to each other and records that belong to two different groups are dissimilar. Each such group is called a cluster and each record belongs to exactly one cluster.¹ Similarity between records is measured computationally by a distance function. A distance function takes two input records and returns a value that is a measure of their similarity. Different applications have different notions of similarity, and no one measure works for all domains.

As an example, consider the schema of the CustomerInfo view:

CustomerInfo(*age*: int, *salary*: real)

We can plot the records in the view on a two-dimensional plane as shown in Figure 26.12. The two coordinates of a record are the values of the record's *salary* and *age* fields. We can visually identify three clusters: Young customers who have low salaries, young customers with high salaries, and older customers with high salaries.

Usually, the output of a clustering algorithm consists of a summarized representation of each cluster. The type of summarized representation depends strongly on the type and shape of clusters the algorithm computes. For example, assume that we have spherical clusters as in the example shown in Figure 26.12. We can summarize each cluster by its *center* (often also called the *mean*) and its *radius*, which are defined as follows. Given a collection of records r_1, \dots, r_n , their center C and radius R are defined as follows:

$$C = \frac{1}{n} \sum_{i=1}^n r_i, \text{ and } R = \sqrt{\frac{\sum_{i=1}^n (r_i - C)^2}{n}}$$

¹There are clustering algorithms that allow overlapping clusters, where a record could belong to several clusters.

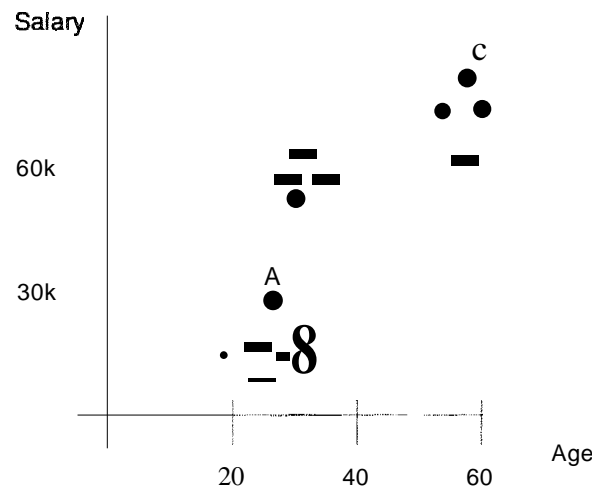


Figure 26.12 Records in CustomerInfo

There are two types of clustering algorithms. A **partitional** clustering algorithm partitions the data into k groups such that some criterion that evaluates the clustering quality is optimized. The number of clusters k is a parameter whose value is specified by the user. A **hierarchical** clustering algorithm generates a sequence of partitions of the records. Starting with a partition in which each cluster consists of one single record, the algorithm merges two partitions in each step until only one single partition remains in the end.

26.5.1 A Clustering Algorithm

Clustering is a very old problem, and numerous algorithms have been developed to cluster a collection of records. Traditionally, the number of records in the input database was assumed to be relatively small and the complete database was assumed to fit into main memory. In this section, we describe a clustering algorithm called BIRCH that handles very large databases. The design of BIRCH reflects the following two assumptions:

- The number of records is potentially very large, and therefore we want to make only one scan over the database.
- Only a limited amount of main memory is available.

A user can set two parameters to control the BIRCH algorithm. The first is a threshold on the amount of main memory available. This main memory threshold translates into a maximum number of cluster summaries k that can be maintained in memory. The second parameter ϵ is an initial threshold for the radius of any cluster. The value of ϵ is an upper bound on the radius of any cluster and controls the number of clusters that the algorithm discovers. If ϵ is small, we discover many small clusters; if ϵ is large, we discover very few

clusters, each of which is relatively large. We say that a cluster is compact if its radius is smaller than ϵ .

BIRCH always maintains k or fewer cluster summaries (C_i, R_i) in main memory, where C_i is the center of cluster i and R_i is the radius of cluster i . The algorithm always maintains compact clusters; that is, the radius of each cluster is less than ϵ . If this invariant cannot be maintained with the given amount of main memory, ϵ is increased as described next.

The algorithm reads records from the database sequentially and processes them as follows:

1. Compute the distance between record r and each of the existing cluster centers. Let i be the cluster index such that the distance between r and C_i is the smallest.
2. Compute the value of the new radius R'_i of the i th cluster under the assumption that r is inserted into it. If $R'_i \leq \epsilon$, then the i th cluster remains compact, and we assign r to the i th cluster by updating its center and setting its radius to R'_i . If $R'_i > \epsilon$, then the i th cluster would no longer be compact if we insert r into it. Therefore, we start a new cluster containing only the record r .

The second step presents a problem if we already have the maximum number of cluster summaries, k . If we now read a record that requires us to create a new cluster, we lack the main memory required to hold its summary. In this case, we increase the radius threshold ϵ —using some heuristic to determine the increase—in order to *merge* existing clusters: An increase of ϵ has two consequences. First, existing clusters can accommodate more records, since their maximum radius has increased. Second, it might be possible to merge existing clusters such that the resulting cluster is still compact. Thus, an increase in ϵ usually reduces the number of existing clusters.

The complete BIRCH algorithm uses a balanced in-memory tree, which is similar to a B+ tree in structure, to quickly identify the closest cluster center for a new record. A description of this data structure is beyond the scope of our discussion.

26.6 SIMILARITY SEARCH OVER SEQUENCES

A lot of information stored in databases consists of sequences. In this section, we introduce the problem of similarity search over a collection of sequences. Our query model is very simple: We assume that the user specifies a query sequence and wants to retrieve all data sequences that are similar to the

Commercial Data Mining Systems: There are a number of data mining products on the market today, such as SAS Enterprise Miner, SPSS Clementine, CART from Salford Systems, Megaputer PolyAnalyst, ANGOSS KnowledgeStudio. We highlight two that have strong database ties.

IBM's Intelligent Miner offers a wide range of algorithms, including association rules, regression, classification, and clustering. The emphasis of Intelligent Miner is on scalability—the product contains versions of all algorithms for parallel computers and is tightly integrated with IBM's DB2 database system. DB2's object-relational capabilities can be used to define the data mining classes of SQL/MM. Of course, other data mining vendors can use these capabilities to add their own data mining models and algorithms to DB2.

Microsoft's SQL Server 2000 has a component called the Analysis Server that makes it possible to create, apply, and manage data mining models within the DBMS. (SQL Server's OLAP capabilities are also packaged in the Analysis Server component.) The basic approach taken is to represent a mining model as a table; clustering and decision tree models are currently supported. The table conceptually has one row for each possible combination of input (predictor) attribute values. The model is created using a statement analogous to SQL's CREATE TABLE that describes the input on which the model is to be trained and the algorithm to use in constructing the model. An interesting feature is that the input table can be defined, using a specialized view mechanism, to be a *nested table*. For example, we can define an input table with one row per customer, where one of the fields is a nested table that describes the customer's purchases. The SQL/MM extensions for data mining do not provide this capability because SQL:1999 does not currently support nested tables (Section 23.2.1). Several properties of attributes, such as whether they are discrete or continuous, can also be specified.

A model is trained by inserting rows into it, using the INSERT command. It is applied to a new dataset to make predictions using a new kind of join called PREDICTION JOIN; in principle, each input tuple is matched with the corresponding tuple in the mining model to determine the value of the predicted attribute. Thus, end users can create, train, and apply decision trees and clustering using extended SQL. There are also commands to browse models. Unfortunately, users cannot add new models or new algorithms for models, a capability that is supported in the SQL/MIME proposal.

query sequence. Similarity search is different from 'normal' queries in that we are interested not only in sequences that match the query sequence exactly but also those that differ only slightly from the query sequence.

We begin by describing sequences and similarity between sequences. A data sequence X is a series of numbers $X = \langle x_1, \dots, x_k \rangle$. Sometimes X is also called a time series. We call k the length of the sequence. A subsequence $Z = \langle z_1, \dots, z_j \rangle$ is obtained from another sequence $X = \langle x_1, \dots, x_k \rangle$ by deleting numbers from the front and back of the sequence X . Formally, Z is a subsequence of X if $z_1 = x_i, z_2 = x_{i+1}, \dots, z_j = x_{i+j-1}$ for some $i \in \{1, \dots, k-j+1\}$. Given two sequences $X = \langle x_1, \dots, x_k \rangle$ and $Y = \langle y_1, \dots, y_l \rangle$, we can define the Euclidean **Distance** as the distance between the two sequences as follows:

$$\|X - Y\| = \sum_{i=1}^k (x_i - y_i)^2$$

Given a user-specified query sequence and a threshold parameter ϵ , our goal is to retrieve all data sequences that are within ϵ -distance of the query sequence.

Similarity queries over sequences can be classified into two types.

- **Complete Sequence Matching:** The query sequence and the sequences in the database have the same length. Given a user-specified threshold parameter ϵ , our goal is to retrieve all sequences in the database that are within ϵ -distance to the query sequence.
- **Subsequence Matching:** The query sequence is shorter than the sequences in the database. In this case, we want to find all subsequences of sequences in the database such that the subsequence is within distance ϵ of the query sequence. We do not discuss subsequence matching.

26.6.1 An Algorithm to Find Similar Sequences

Given a collection of data sequences, a query sequence, and a distance threshold ϵ , how can we efficiently find all sequences within ϵ -distance of the query sequence?

One possibility is to scan the database, retrieve each data sequence, and compute its distance to the query sequence. While this algorithm has the merit of being simple, it always retrieves every data sequence.

Because we consider the complete sequence matching problem, all data sequences and the query sequence have the same length. We can think of this similarity search as a high-dimensional indexing problem. Each data sequence

and the query sequence can be represented as a point in a k -dimensional space. Therefore, if we insert all data sequences into a multidimensional index, we can retrieve data sequences that exactly match the query sequence by querying the index. But since we want to retrieve not only data sequences that match the query exactly but also all sequences within ϵ -distance of the query sequence, we do not use a point query as defined by the query sequence. Instead, we query the index with a hyper-rectangle that has side-length 2ϵ and the query sequence as center, and we retrieve all sequences that fall within this hyper-rectangle. We then discard sequences that are actually further than ϵ away from the query sequence.

Using the index allows us to greatly reduce the number of sequences we consider and decreases the time to evaluate the similarity query significantly. The bibliographic notes at the end of the chapter provide pointers to further improvements.

26.7 INCREMENTAL MINING AND DATA STREAMS

Real-life data is not static, but is constantly evolving through additions or deletions of records. In some applications, such as network monitoring, data arrives in such high-speed streams that it is infeasible to store the data for offline analysis. We describe both evolving and streaming data in terms of a framework called block evolution. In block evolution, the input dataset to the data mining process is not static but periodically updated with a new block of tuples, for example, every day at midnight or in a continuous stream. A block is a set of tuples added simultaneously to the database. For large blocks, this model captures common practice in many of today's data warehouse installations, where updates from operational databases are batched together and performed in a block update. For small blocks of data—at the extreme, each block consists of a single record—this model captures streaming data.

In the block evolution model, the database consists of a (conceptually infinite) sequence of data blocks D_1, D_2, \dots that arrive at times $1, 2, \dots$, where each block D_i consists of a set of records.² We call i the *block identifier* of block D_i . Therefore, at any time t , the database consists of a finite sequence of blocks of data $\langle D_1, \dots, D_t \rangle$ that arrived at times $\{1, 2, \dots, t\}$. The database at time t , which we denote by $D[1, t]$, is the union of the database at time $t - 1$ and the block that arrives at time t , D_t .

For evolving data, two classes of problems are of particular interest: model maintenance and change detection. The goal of model maintenance is to

²In general, a block specifies records to change or delete, in addition to records to insert. We only consider inserts.

maintain a data mining model under insertion and deletions of blocks of data. To incrementally compute the data mining model at time t , which we denote by $M(D[1, t])$, we must consider only $M(D[1, t - 1])$ and D_t ; we cannot consider the data that arrived prior to time t . Further, a data analyst might specify time-dependent subsets of $D[1, t]$, such as a window of interest (e.g., all the data seen thus far or last week's data). More general selections are also possible, for example, all weekend data over the past year. Given such selections, we must incrementally compute the model on the appropriate subset of $D[1, t]$ by considering only J_t and the model on the appropriate subset of $D[1, t - 1]$. 'Almost' incremental algorithms that occasionally examine older data might be acceptable in warehouse applications, where incrementality is motivated by efficiency considerations and older data is available to us if necessary. This option is not available for high-speed data streams, where older data may not be available at all.

The goal of change detection is to quantify the difference, in terms of their data characteristics, between two sets of data and determine whether the change is meaningful (i.e., statistically significant). In particular, we must quantify the difference between the models of the data as it existed at some time t_1 and the evolved version at a subsequent time t_2 ; that is, we must quantify the difference between $M(D[1, t_1])$ and $M(D[1, t_2])$. We can also measure changes with respect to selected subsets of data. Several natural variants of the problem exist; for example, the difference between $M(D[1, t - 1])$ and $M(D_t)$ indicates whether the latest block differs substantially from previously existing data. In the rest of this chapter, we focus on model maintenance and do not discuss change detection.

Incremental model maintenance has received much attention. Since the quality of the data mining model is of utmost importance, incremental model maintenance algorithms have concentrated on computing exactly the same model as computed by running the basic model construction algorithm on the union of old and new data. One widely used scalability technique is localization of changes due to new blocks. For example, for density-based clustering algorithms, the insertion of a new record affects only clusters in the neighborhood of the record, and thus efficient algorithms can *localize* the change to a few clusters and avoid recomputing all clusters. As another example, in decision tree construction, we might be able to show that the split criterion at a node of the tree changes only within acceptably small confidence intervals when records are inserted, if we assume that the underlying distribution of training records is static.

One-pass model construction over data streams has received particular attention, since data arrives and must be processed continuously in several emerging applications. For example, network installations of large Telecom

and Internet service providers have detailed usage information (e.g., call-detail-records, router packet-flow and trace data) from different parts of the underlying network that needs to be continuously analyzed to detect interesting trends. Other examples include webserver logs, streams of transactional data from large retail chains, and financial stock tickers.

When working with high-speed data streams, algorithms must be designed to construct data mining models while looking at the relevant data items *only once and in a fixed order* (determined by the stream-arrival pattern), with a limited amount of main memory. Data-stream computation has given rise to several recent (theoretical and practical) studies of online or one-pass algorithms with bounded memory. Algorithms have been developed for one-pass computation of quantiles and order-statistics, estimation of frequency moments and join sizes, clustering and decision tree construction, estimating correlated aggregates, and computing one-dimensional (i.e., single-attribute) histograms and Haar wavelet decompositions. Next, we discuss one such algorithm, for incremental maintenance of frequent itemsets.

26.7.1 Incremental Maintenance of Frequent Itemsets

Consider the Purchases Relation shown in Figure 26.1 and assume that the minimum support threshold is 60%. It can be easily seen that the set of frequent itemsets of size 1 consists of $\{pen\}$, $\{ink\}$, and $\{milk\}$ with supports of 100%, 75%, and 75%, respectively. The set of frequent itemsets of size 2 consists of $\{pen, ink\}$ and $\{pen, milk\}$, both with supports of 75%. The Purchases relation is our first block of data. Our goal is to develop an algorithm that maintains the set of frequent itemsets under insertion of new blocks of data.

As a first example, let us consider the addition of the block of data shown in Figure 26.13 to our original database (Figure 26.1). Under this addition, the set of frequent itemsets does not change, although their support values do: $\{pen\}$, $\{ink\}$, and $\{milk\}$ now have support values of 100%, 60%, and 60%, respectively, and $\{pen, ink\}$ and $\{pen, milk\}$ now have 60% support. Note that we could detect this case of 'no change' simply by maintaining the number of market baskets in which each itemset occurred. In this example, we update the (absolute) support of itemset $\{pen\}$ by 1.

<i>transid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
115	201	7/1/99	pen	2

Figure 26.13 The Purchases Relation Block 2

<i>transid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
115	<u>201</u>	7/1/99	water	1
115	201	7/1/99	llilk	1

Figure 26.14 The Purchases Relation Block 2a

In general, the set of frequent itemsets may change. As an example, consider the addition of the block shown in Figure 26.14 to the original database shown in Figure 26.1. We see a transaction containing the item *water*, but we do not know the support of the itemset $\{\textit{water}\}$, since *water* was not above the minimum support in our original database. A simple solution in this case is to make an additional scan over the original database and compute the support of the itemset $\{\textit{water}\}$. But can we do better? Another immediate solution is to keep counters for *all* possible itemsets, but the number of all possible itemsets is exponential in the number of items—and most of these counters would be 0 anyway. Can we design an intelligent strategy that tells us *which* counters to maintain?

We introduce the notion of the *negative border* of a set of itemsets to help decide which counters to keep. The negative border of a set of frequent itemsets consists of all itemsets X such that X itself is not frequent, but all subsets of X are frequent. For example, in the case of the database shown in Figure 26.1, the following itemsets make up the negative border: $\{\textit{juice}\}$, $\{\textit{water}\}$, and $\{\textit{ink}, \textit{milk}\}$. Now we can design a more efficient algorithm for maintaining frequent itemsets by keeping counters for all currently frequent itemsets *and* all itemsets currently in the negative border. (Only if an itemset in the negative border becomes frequent do we need to read the original dataset again, to find the support for new candidate itemsets that might be frequent.)

We illustrate this point through the following two examples. If we add Block 2a shown in Figure 26.14 to the original database shown in Figure 26.1, we increase the support of the frequent itemset $\{\textit{milk}\}$ by one, and we increase the support of the itemset $\{\textit{water}\}$, which is in the negative border, by one as well. But since no itemset in the negative border became frequent, we do not have to re-scan the original database.

In contrast, consider the addition of Block 2b shown in Figure 26.15 to the original database shown in Figure 26.1. In this case, the itemset $\{\textit{juice}\}$, which was originally in the negative border, becomes frequent with a support of 60%. This means that now the following itemsets of size two enter the negative border: $\{\textit{juice}, \textit{pen}\}$, $\{\textit{juice}, \textit{ink}\}$, and $\{\textit{juice}, \textit{milk}\}$. (We know that $\{\textit{juice}, \textit{water}\}$ cannot be frequent since the itemset $\{\textit{water}\}$ is not frequent.)

115	201	7/1/99	juice	2
115	201	7/1/99	water	2

Figure 26.15 The Purchases Relation Block 2b

26.8 ADDITIONAL DATA MINING TASKS

We focused on the problem of discovering patterns from a database, but there are several other equally important data mining tasks. We now discuss some of these briefly. The bibliographic references at the end of the chapter provide many pointers for further study.

- **Dataset and Feature Selection:** It is often important to select the ‘right’ dataset to mine. Dataset selection is the process of finding which datasets to mine. Feature selection is the process of deciding which attributes to include in the mining process.
- **Sampling:** One way to explore a large dataset is to obtain one or more *samples* and analyze them. The advantage of sampling is that we can carry out detailed analysis on a sample that would be infeasible on the entire dataset, for very large datasets. The disadvantage of sampling is that obtaining a representative sample for a given task is difficult; we might miss important trends or patterns because they are not reflected in the sample. Current database systems also provide poor support for efficiently obtaining samples. Improving database support for obtaining samples with various desirable statistical properties is relatively straightforward and likely to be available in future DBMSs. Applying sampling for data mining is an area for further research.
- **Visualization:** Visualization techniques can significantly assist in understanding complex datasets and detecting interesting patterns, and the importance of visualization in data mining is widely recognized.

26.9 REVIEW QUESTIONS

Answers to the review questions can be found in the listed sections.

- What is the role of data mining in the KDD process? (Section 26.1)
- What is the a priori property? Describe an algorithm for finding frequent itemsets. (Section 26.2.1)

- How are iceberg queries related to frequent itemsets? (Section 26.2.2)
- Give the definition of an *association rule*. What is the difference between support and confidence of a rule? (Section 26.3.1)
- Can you explain extensions of association rules to ISA hierarchies? What other extensions of association rules are you familiar with? (Sections 26.3.3 and 26.3.4)
- What is a sequential pattern? How can we compute sequential patterns? (Section 26.3.5)
- Can we use association rules for prediction? (Section 26.3.6)
- What is the difference between Bayesian Networks and association rules? (Section 26.3.7)
- Can you give examples of classification and regression rules? How is support and confidence for such rules defined? (Section 26.3.8)
- What are the components of a decision tree? How are decision trees constructed? (Sections 26.4.1 and 26.4.2)
- What is a cluster? What information do we usually output for a cluster? (Section 26.5)
- How can we define the distance between two sequences? Describe an algorithm to find all sequences similar to a query sequence. (Section 26.6)
- Describe the block evolution model and define the problems of incremental model maintenance and change detection. What is the added challenge in mining data streams? (Section 26.7)
- Describe an incremental algorithm for computing frequent itemsets. (Section 26.7.1)
- Give examples of other tasks related to data mining. (Section 26.8)

EXERCISES

Exercise 26.1 Briefly answer the following questions:

1. Define *support* and *confidence* for an association rule.
2. Explain why association rules cannot be used directly for prediction, without further analysis or domain knowledge.
3. What are the differences between *association rules*, *classification rules*, and *regression rules*?
4. What is the difference between *classification* and *clustering*?

<i>pid</i>	<i>custid</i>	<i>date</i>	<i>item</i>	<i>qty</i>
111	201	5/1/2002	ink	1
111	201	5/1/2002	lnilk	2
111	201	5/1/2002	JuIce	1
112	105	6/3/2002	pen	1
112	105	6/3/2002	ink	1
112	105	6/3/2002	water	1
113	106	5/10/2002	pen	1
113	106	5/10/2002	water	2
113	106	5/10/2002	milk	1
114	201	6/1/2002	pen	2
114	201	6/1/2002	ink	2
114	201	6/1/2002	JuIce	4
114	201	6/1/2002	water	1
114	201	6/1/2002	ruilk	1

Figure 26.16 The Purchases2 Relation

5. What is the role of information visualization in data mining?
6. Give examples of queries over a database of stock price quotes, stored as sequences, one per stock, that cannot be expressed in SQL.

Exercise 26.2 Consider the Purchases table shown in Figure 26.1.

1. Simulate the algorithm for finding frequent itemsets on the table in Figure 26.1 with $\text{minsup}=90$ percent, and then find association rules with $\text{minconf}=90$ percent.
2. Can you modify the table so that the same frequent itemsets are obtained with $\text{minsup}=90$ percent as with $\text{minsup}=70$ percent on the table shown in Figure 26.1?
3. Simulate the algorithm for finding frequent itemsets on the table in Figure 26.1 with $\text{minsup}=10$ percent and then find association rules with $\text{minconf}=90$ percent.
4. Can you modify the table so that the same frequent itemsets are obtained with $\text{minsup}=10$ percent as with $\text{minsup}=70$ percent on the table shown in Figure 26.1?

Exercise 26.3 Assume we are given a dataset D of market baskets and have computed the set of frequent itemsets \mathcal{X} in D for a given support threshold minsup . Assume that we would like to add another dataset D' to D , and maintain the set of frequent itemsets with support threshold minsup in $D \cup D'$. Consider the following algorithm for incremental maintenance of a set of frequent itemsets:

1. We run the *a priori* algorithm on D' and find all frequent itemsets in D' and their support. The result is a set of itemsets \mathcal{X}' . We also compute the support of all itemsets $X \in \mathcal{X}$ in D' .
2. We then make a scan over D to compute the support of all itemsets in \mathcal{X}' .

Answer the following questions about the algorithm:

- The last step of the algorithm is missing; that is, what should the algorithm output?
- Is this algorithm more efficient than the algorithm described in Section 26.7.1?

Exercise 26.4 Consider the Purchases2 table shown in Figure 26.16.

- List all itemsets in the negative border of the dataset.
- List all frequent itemsets for a support threshold of 50%.
- Give an example of a database in which the addition of this database does not change the negative border.
- Give an example of a database in which the addition of this database would change the negative border.

Exercise 26.5 Consider the Purchases table shown in Figure 26.1. Find all (generalized) association rules that indicate the likelihood of items being purchased on the same date by the same customer, with *minsup* set to 10% and *minconj* set to 70%.

Exercise 26.6 Let us develop a new algorithm for the computation of all large itemsets. Assume that we are given a relation D similar to the Purchases table shown in Figure 26.1. We partition the table horizontally into k parts D_1, \dots, D_k .

1. Show that, if itemset X is frequent in D , then it is frequent in at least one of the k parts.
2. Use this observation to develop an algorithm that computes all frequent itemsets in two scans over D . (Hint: In the first scan, compute the locally frequent itemsets for each part D_i , $i \in \{1, \dots, k\}$.)
3. Illustrate your algorithm using the Purchases table shown in Figure 26.1. The first partition consists of the two transactions with *transid* 111 and 112, the second partition consists of the two transactions with *transid* 113 and 114. Assume that the minimum support is 70 percent.

Exercise 26.7 Consider the Purchases table shown in Figure 26.1. Find all sequential patterns with *minsup* set to 60%. (The text only sketches the algorithm for discovering sequential patterns, so use brute force or read one of the references for a complete algorithm.)

Exercise 26.8 Consider the SubscriberInfo Relation shown in Figure 26.17. It contains information about the marketing campaign of the *DB Aficionado* magazine. The first two columns show the age and salary of a potential customer and the *subscription* column shows whether the person subscribes to the magazine. We want to use this data to construct a decision tree that helps predict whether a person will subscribe to the magazine.

1. Construct the AVC-group of the root node of the tree.
2. Assume that the splitting predicate at the root node is $age \leq 50$. Construct the AVC-groups of the two children nodes of the root node.

Exercise 26.9 Assume you are given the following set of six records: $(7, 55)$, $(21, 202)$, $(25, 220)$, $(12, 73)$, $(8, 61)$, and $(22, 249)$.

1. Assuming that all six records belong to a single cluster, compute its center and radius.
2. Assume that the first three records belong to one cluster and the second three records belong to a different cluster. Compute the center and radius of the two clusters.
3. Which of the two clusterings is 'better' in your opinion and why?

Exercise 26.10 Assume you are given the three sequences $(1, 3, 4)$, $(2, 3, 2)$, $(3, 3, 7)$. Compute the Euclidean Distance between all pairs of sequences.

<u>age</u>	<u>salary</u>	<u>subscription</u>
37	45k	No
39	70k	Yes
56	50k	Yes
52	43k	Yes
35	90k	Yes
32	54k	No
40	58k	No
55	85k	Yes
43	68k	Yes

Figure 26.17 The SubscriberInfo Relation

BIBLIOGRAPHIC NOTES

Discovering useful knowledge from a large database is more than just applying a collection of data mining algorithms, and the point of view that it is an iterative process guided by an analyst is stressed in [265] and [666]. Work on exploratory data analysis in statistics, for example [745], and on machine learning and knowledge discovery in artificial intelligence was a precursor to the current focus on data mining; the added emphasis on large volumes of data is the important new element. Good recent surveys of data mining algorithms include [267, 397, 507]. [266] contains additional surveys and articles on many aspects of data mining and knowledge discovery, including a tutorial on Bayesian networks [371]. The book by Piatetsky-Shapiro and Frawley [595] contains an interesting collection of data mining papers. The annual SIGKDD conference, run by the ACM special interest group in knowledge discovery in databases, is a good resource for readers interested in current research in data mining [25, 162, 268, 372, 613, 691], as is the *Journal of Knowledge Discovery and Data Mining*. [363, 370, 511, 781] are good, in-depth textbooks on data mining.

The problem of mining association rules was introduced by Agrawal, Imielinski, and Swami [20]. Many efficient algorithms have been proposed for the computation of large itemsets, including [21, 117, 364, 683, 738, 786].

Iceberg queries have been introduced by Fang et al. [264]. There is also a large body of research on generalized forms of association rules; for example, [700, 701, 703]. The problem of finding maximal frequent itemsets has also received significant attention [13, 67, 126, 346, 347, 479, 787]. Algorithms for mining association rules with constraints are considered in [68, 462, 563, 590, 591, 703].

Parallel algorithms are described in [23] and [655]. Recent papers on parallel data mining can be found in [788], and work on distributed data mining can be found in [417].

[291] presents an algorithm for discovering association rules over a continuous numeric attribute; association rules over numeric attributes are also discussed in [783]. The general form of association rules, in which attributes other than the transaction id are grouped is developed in [529]. Association rules over items in a hierarchy are discussed in [361, 700]. Further

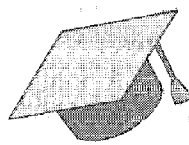
extensions and generalization of association rules are proposed in [67, 115, 563]. Integration of mining for frequent itemsets into database systems has been addressed in [654, 743]. The problem of mining sequential patterns is discussed in [24], and further algorithms for mining sequential patterns can be found in [510, 702].

General introductions to classification and regression rules can be found in [362, 532]. The classic reference for decision and regression tree construction is the CART book by Breilnan, Friedmau, Olsheu, and Stone [111]. A machine learning perspective of decision tree construction is given by Quinlan [603]. Recently, several scalable algorithms for decision tree construction have been developed [309, 311, 521, 619, 674].

The clustering problem has been studied for decades in several disciplines. Sample textbooks include [232, 407, 418]. Scalable clustering algorithms include CLARANS [562], DBSCAN [249, 250], BIRCH [798], and CURE [344]. Bradley, Fayyad, and Reina address the problem of scaling the K-Means clustering algorithm to large databases [108, 109]. The problem of finding clusters in subsets of the fields is addressed in [19]. Ganti et al. examine the problem of clustering data in arbitrary metric spaces [302]. Algorithms for clustering categorical data include STIRR [315] and CACTUS [301]. [651] is a clustering algorithm for spatial data.

Finding similar sequences from a large database of sequences is discussed in [22, 262, 446, 606, 680].

Work on incremental maintenance of association rules is considered in [174, 175, 736]. Ester et al. describe how to maintain clusters incrementally [248], and Hidber describes how to maintain large itemsets incrementally [378]. There has also been recent work on mining data streams, such as the construction of decision trees over data streams [228, 309, 393] and clustering data streams [343, 568]. A general framework for mining evolving data is presented in [299]. A framework for measuring change in data characteristics is proposed in [300].



27

INFORMATION RETRIEVAL AND XML DATA

- How are DBMSs evolving in response to the growing alllounts of text data?
- What is the vector space rmodel and how does it support text search?
- How are text collections indexed?
- Cornpared to IR systemls, what is new in Web search?
- How is XML data different from plain text and relational tables?
- What are the main features of XQuery?
- What are the irnplementation challenges posed by XML data?
- Key concepts: information retrieval, boolean and ranked queries; relevance, precision, recall; vector space model, TF/IDF ternn weight-ing, document similarity; inverted index, signature file; Web crawler, hubs and authorities, Pigeon Rank of a webpage; sernistructured data llmodel, XML; XQuery, path expressions, FLWR queries; XML storage and indexing

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A *memex* is a device in which an individual stores all his books, records, and cornrnunications, and which is rnechanized so that it rnay be consulted with exceeding speed and flexibility.

—Vannevar Bush, *As We May Think*, 1945

The field of information retrieval (IR) has studied the problem of searching collections of text documents since the 1950s and developed largely independently of database systems. The proliferation of text documents on the Web made document search an everyday operation for most people and led to renewed research on the topic.

The database field's desire to expand the kinds of data that can be managed in a DBMS is well-established and reflected in developments like object-relational extensions (Chapter 23). Documents on the Web represent one of the most rapidly growing sources of data, and the challenge of managing such documents in a DBMS has naturally become a focal point for database research.

The Web, therefore, brought the two fields of database management systems and information retrieval closer together than ever before, and, as we will see, XML sits squarely in the middle ground between them. We introduce IR systems as well as a data model and query language for XML data and discuss the relationship with (object-)relational database systems.

In this chapter, we present an overview of information retrieval, Web search, and the emerging XML data model and query language standards. We begin in Section 27.1 with a discussion of how these text-oriented trends fit within the context of current object-relational database systems. We introduce information retrieval concepts in Section 27.2 and discuss specialized indexing techniques for text in Section 27.3. We discuss Web search engines in Section 27.4. In Section 27.5, we briefly outline current trends in extending database systems to support text data and identify some of the important issues involved. In Section 27.6, we present the XML data model, building on the XML concepts introduced in Chapter 7. We describe the XQuery language in Section 27.7. In Section 27.8, we consider efficient evaluation of XQuery queries.

27.1 COLLIDING WORLDS: DATABASES, IR, AND XML

The Web is the most widely used document collection today, and search on the Web differs from traditional IR-style document retrieval in important ways. First, there is great emphasis on scalability to very large document collections. IR systems typically dealt with tens of thousands of documents, whereas the Web contains billions of pages.

Second, the Web has significantly changed how document collections are created and used. Traditionally, IR systems were aimed at professionals like librarians and legal researchers, who were trained in using sophisticated retrieval engines. Documents were carefully prepared, and documents in a given collection were typically on related topics. On the Web, documents are created by an infinite

variety of individuals for equally many purposes, and reflect this diversity in size and content. Searches are carried out by ordinary people with no training in using retrieval software.

The emergence of XML has added a third interesting dimension to text search: Every document can now be marked up to reflect additional information of interest, such as authorship, source, and even details about the intrinsic content. This has changed the nature of a “document” from free text to textual objects with associated fields containing metadata (data about data) or descriptive information. Links to other documents are a particularly important kind of metadata, and they can have great value in searching document collections on the Web.

The Web also changed the notion of what constitutes a document. Documents on the Web may be multimedia objects such as images or video clips, with text appearing only in descriptive tags. We must be able to manage such heterogeneous data collections and support searches over them.

Database management systems traditionally dealt with simple tabular data. In recent years, object-relational database systems (ORDBMSs) were designed to support complex data types. Images, videos, and textual objects have been explicitly mentioned as examples of the data types ORDBMSs are intended to support. Nonetheless, current database systems have a long way to go before they can support such complex data types satisfactorily. In the context of text and XML data, challenges include efficient support for searches over textual content and support for searches that exploit the loose structure of XML data.

27.1.1 DBMS versus IR Systems

Database and IR systems have the common objective of supporting searches over collections of data. However, many important differences have influenced their development.

- **Searches versus Queries:** IR systems are designed to support a specialized class of queries that we also call searches. Searches are specified in terms of a few search terms, and the underlying data is usually a collection of unstructured text documents. In addition, an important feature of IR searches is that search results may be ranked, or ordered, in terms of how ‘well’ the search results match the search terms. In contrast, database systems support a very general class of queries, and the underlying data is rigidly structured. Unlike IR systems, database systems have traditionally returned unranked sets of results. (Even the recent SQL/OLAP extensions that support early results and searches over ordered data (see Chapter 25)

do not order results in terms of how well they match the query. Relational queries are *precise* in that a row is either in the answer or it is not; there is no notion of 'how well a row matches' the query.) In other words, a relational query only assigns two ranks to a row, indicating 'whether the row is in the answer or not.

- **Updates and Transactions:** IR systems are optimized for a read-mostly workload and do not support the notion of a transaction. In traditional IR systems, new documents are added to the document collection from time to time, and index structures that speed up searches are periodically rebuilt or updated. Therefore, documents that are highly relevant for a search might exist in the IR system, but not be retrievable yet because of outdated index structures. In contrast, database systems are designed to handle a wide range of workloads, including update-intensive transaction processing workloads.

These differences in design objectives have led, not surprisingly, to very different research emphases and system designs. Research in IR studied ranking functions extensively. For example, among other topics, research in IR investigated how to incorporate feedback from a user's behavior to modify a ranking function and how to apply linguistic processing techniques to improve searches. Database research concentrated on query processing, concurrency control and recovery, and other topics, as covered in this book.

The differences between a DBMS and an IR system from a design and implementation standpoint should become clear as we introduce IR systems in the next few sections.

27.2 INTRODUCTION TO INFORMATION RETRIEVAL

There are two common types of searches, or queries, over text collections: boolean queries and ranked queries. In a **boolean query**, the user specifies an expression constructed using terms and boolean operators (And, Or, Not). For example,

database And (Microsoft Or IBM)

This query asks for all documents that contain the term *database* and in addition, either *Microsoft* or *IBM*.

In a ranked query the user specifies one or more terms, and the result of the query is a list of documents ranked by their relevance to the query. Intuitively, documents at the top of the result list are expected to 'match' the search

docid	Document
1	agent Janles Bond good agent
2	agent rnobile cornputer
3	James Madison Inovie
4	Janles Bond movie

Figure 27.1 A Text Database with Four Records

condition ruore closely, or be 'rnore relevant', than doculnents lower in the result list. While a document that contains *Microsoft* satisfies the search '*Microsoft, IBM,*' a document that also contains *IBM* is considered to be a better match. Similarly, a docunlent that contains several occurrences of *Microsoft* might be a better rnatch than a document that contains a single occurence. Ranking the docurnents that satisfy the boolean search condition is an important aspect of an IR search engine, and we discuss how this is done in Sections 27.2.3 and 27.4.2.

An important extension of ranked queries is to ask for documents that are most relevant to a given natural language sentence. Since a sentence has linguistic structure (e.g., subject-verb-object relationships), it provides more information than just the list of words that it contains. We do not discuss natural language search.

27.2.1 Vector Space Model

We now describe a widely-used franlework for representing docurnents and searching over document collections. Consider the set of all terns that appear in a given collection of documents. We can represent each document as a vector with one entry per ternl. In the shnplest 1'01'111 of docunlent vectors, if tern j appears k times in dOCUlnt i , the document vector for document i contains value k in position j . The document vector for i contains the value 0 in positions corresponding to terns that do not appear in i .

Consider the exaInple collection of four documents shown in Figure 27.1. rre docUluent vector representation is illustrated in Figure 27.2; each row represents a document. This representation of documents as tern vectors is called the vector space model.

docid	agent	Bond	computer	good	James	Madison	mobile	movie
1	2	1	0	1	1	0	0	0
2	1	0	1	0	0	0	1	0
3	0	0	0	0	1	1	0	1
4	0	1	0	0	1	0	0	1

Figure 27.2 Document Vectors for the Example Collection

27.2.2 TFIDF Weighting of Terms

We described the value for a term in a document vector as simply the term frequency (TF), or number of occurrences of that term in the given document. This reflects the intuition that a term which appears often is more important in characterizing the document than a term that appears only once (or a term that does not appear at all).

However, some terms appear very frequently in the document collection, and others are relatively rare. The frequency of terms is empirically observed to follow a Zipfian distribution, as illustrated in Figure 27.3. In this figure, each position on the X-axis corresponds to a term and the Y-axis corresponds to the number of occurrences of the term. Terms are arranged on the X-axis in decreasing order by the number of times they occur (in the document collection as a whole).

As might be expected, it turns out that extremely common terms are not very useful in searches. Examples of such common terms include *a*, *an*, *the* etc. Terms that occur extremely often are called stop words, and documents are pre-processed to eliminate stop words.

Even after eliminating stop words, we have the phenomenon that some words appear much more often than others in the document collection. Consider the words *Linux* and *kernel* in the context of a collection of documents about the Linux operating system. While neither is common enough to be a stop word, *Linux* is likely to appear much more often. Given a search that contains both these keywords, we are likely to get better results if we give more importance to documents that contain *kernel* than documents that contain *Linux*.

We can capture this intuition by refining the document vector representation as follows. The value associated with term j in the document vector for document i , denoted as w_{ij} , is obtained by multiplying the term frequency t_{ij} (the number of times term j appears in document i) by the inverse document frequency (IDF) of term j in the document collection. IDF of a term j is defined as

$\log(1/n_j)$; where N is the total number of documents, and n_j is the number of documents that term j appears in. This effectively increases the weight given to rare terms. As an example, in a collection of 10,000 documents, a term that appears in half the documents has an IDF of 0.3, and a term that occurs in just one document has an IDF of 4.

Length Normalization

Consider a document D . Suppose that we modify it by adding a large number of new terms. Should the weight of a term t that appears in D be the same in the document vectors for D and the modified document? Although the TF-IDF weight for t is indeed the same in the two document vectors, our intuition suggests that the weight should be less in the modified document. Longer documents tend to have more terms, and more occurrences of any given term. Thus, if two documents contain the same number of occurrences of a given term, the importance of the term in characterizing the document also depends on the length of the document.

Several approaches to length normalization have been proposed. Intuitively, all of them reduce the importance given to how often a term occurs as the frequency grows. In traditional IR systems, a popular way to refine the similarity metric is cosine length normalization:

$$w_{ij}^* = \frac{w_{ij}}{\sqrt{\sum_{k=1}^t w_{ik}^2}}$$

In this formula, t is the number of terms in the document collection, w_{ij} is the TF-IDF weight without length normalization, and w_{ij}^* is the length adjusted TF-IDF weight.

Terms that occur frequently in a document are particularly problematic on the Web because webpages are often deliberately modified by adding many copies of certain words... for example, sale, free, sex... to increase the likelihood of their being returned in response to queries. For this reason, Web search engines typically normalize for length by imposing a maximum value (usually 2 or 3) for term frequencies.

27.2.3 Ranking Document Similarity

We now consider how the vector space representation allows us to rank documents in the result of a ranked query. A key observation is that a ranked query can itself be thought of as a document, since it is just a collection of terms. This allows us to use document similarity as the basis for ranking query

results--the document that is most similar to the query is ranked highest, and the one that is least similar is ranked lowest.

If a total of t terms appear in the collection of documents (t is 8 in the example shown in Figure 27.2), we can visualize document vectors in a t -dimensional space in which each axis is labeled with a term. This is illustrated in Figure 27.4, for a two-dimensional space. The figure shows document vectors for two documents, D_1 and D_2 , as well as a query Q .

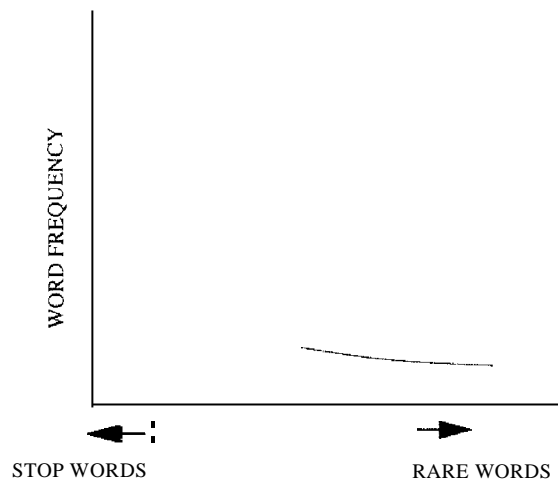


Figure 27.3 Zipfian Distribution of Term Frequencies

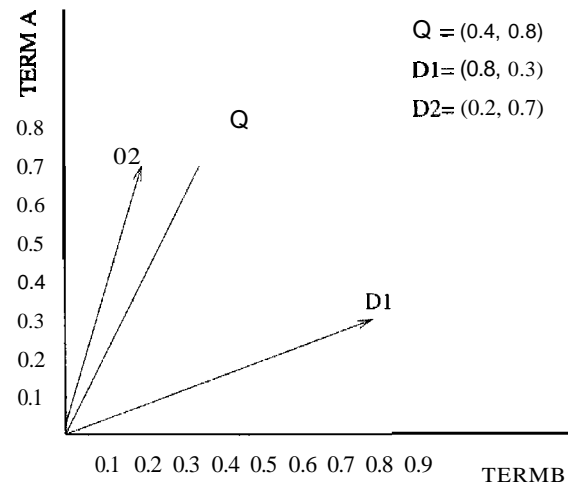


Figure 27.4 Document Similarity

The traditional measure of closeness between two vectors, their *dot product*, is used as a measure of document similarity. The similarity of query Q to a document D_i is illustrated by their dot product:

$$\text{sim}(Q, D_i) = \sum_{j=1}^t q_j^* \cdot w_{ij}^*$$

In the example shown in Figure 27.4, $\text{sim}(Q, D_1) = (0.4 * 0.8) + (0.8 * 0.3) = 0.56$, and $\text{sim}(Q, D_2) = (0.4 * 0.2) + (0.8 * 0.7) = 0.64$. Accordingly, D_2 is ranked higher than D_1 in the search result.

In the context of the Web, document similarity is one of several measures that can be used to rank results, but should not be used exclusively. First, it is questionable whether users want documents that are similar to the query (which typically consists of one or two words) or documents that contain useful information related to the query terms. Intuitively, we want to give importance to the *quality* of a Web page while ranking it, in addition to reflecting the similarity of the page to a given query. Links between pages provide valuable

additional information that can be used to obtain high-quality results. We discuss this issue in Section 27.4.2.

27.2.4 Measuring Success: Precision and Recall

Two criteria are commonly used to evaluate information retrieval systems. Precision is the percentage of retrieved documents that are relevant to the query. Recall is the percentage of relevant documents in the database that are retrieved in response to a query.

Retrieving all documents in response to a query trivially guarantees perfect recall, but results in very poor precision. The challenge is to achieve good recall together with high precision.

In the context of search over the Web, the size of the underlying collection is on the order of billions of documents. Given this, it is questionable whether the traditional measure of recall is very useful. Since users typically don't look beyond the first screen of results, the quality of a Web search engine is largely determined by the results shown on the first page. The following adapted definitions of precision and recall might be more appropriate for Web search engines:

- **Web Search Precision:** The percentage of results on the first page that are relevant to the query.
- **Web Search Recall:** the fraction N/M , expressed as a percentage, where M is the number of results displayed on the front page, and of the M most relevant documents, N is the number displayed on the front page.

27.3 INDEXING FOR TEXT SEARCH

In this section, we introduce two indexing techniques that support the evaluation of boolean and ranked queries. The *'inverted index'* structure discussed in Section 27.3.1 is widely used due to its simplicity and good performance. Its main disadvantage is that it imposes a significant space overhead: The size can be up to 300 percent the size of the original file. The *signature file* index discussed in Section 27.3.2 has a small space overhead and offers a quick filter that eliminates most nonqualifying documents. However, it does not scale as well to larger database sizes because the index has to be sequentially scanned.

Before a document is indexed, it is typically pre-processed to eliminate stop words. Since the size of the indexes is very sensitive to the number of terms in the document collection, eliminating stop words can greatly reduce index

size. IR systems also do certain other kinds of pre-processing. For instance, they apply stemming to reduce related terms to a canonical form. This step also reduces the number of terms to be indexed, but equally importantly, it allows us to retrieve documents that may not contain the exact query term but contain some variant. As an example, the terms *run*, *running*, and *runner* all stem to *run*. The term *run* is indexed, and every occurrence of a variant of this term is treated as an occurrence of *run*. A query that specifies *runner* finds documents that contain any word that stems to *run*.

27.3.1 Inverted Indexes

An inverted index is a data structure that enables fast retrieval of all documents that contain a query term. For each term, the index maintains a list (called the inverted list) of entries describing occurrences of the term, with one entry per document that contains the term.

Consider the inverted index for our running example shown in Figure 27.5. The term 'Janes' has an inverted list with one entry each for documents 1, 3, and 4; the term 'agent' has entries for documents 1 and 2.

The entry for document d in the inverted list for term t contains details about the occurrences of term t in document d . In Figure 27.5, this information consists of a list of locations within the document that contain term t . Thus, the entry for document 1 in the inverted list for term 'agent' lists the locations 1 and 5, since 'agent' is the first and fifth word of document 1. In general, we can store additional information about each occurrence (e.g., in an HTML document, is the occurrence in the TITLE tag?) in the inverted list. We can also store the length of the document if this is used for length normalization (see below).

The collection of inverted lists is called the postings file. Inverted lists can be very large for large document collections. In fact, Web search engines typically store each inverted list on a separate page, and most lists span multiple pages (and if so, are maintained as a linked list of pages). In order to quickly find the inverted list for a query term, all possible query terms are organized in a second index structure such as a B+ tree or a hash index.

The second index, called the lexicon, is much smaller than the postings file since it only contains one entry per term, and further, only contains entries for the set of terms that are retained after eliminating stop words, and applying stemming rules. An entry consists of the term, some summary information about its inverted list, and the address (on disk) of the inverted list. In Figure 27.5, the summary information consists of the number of entries in the inverted

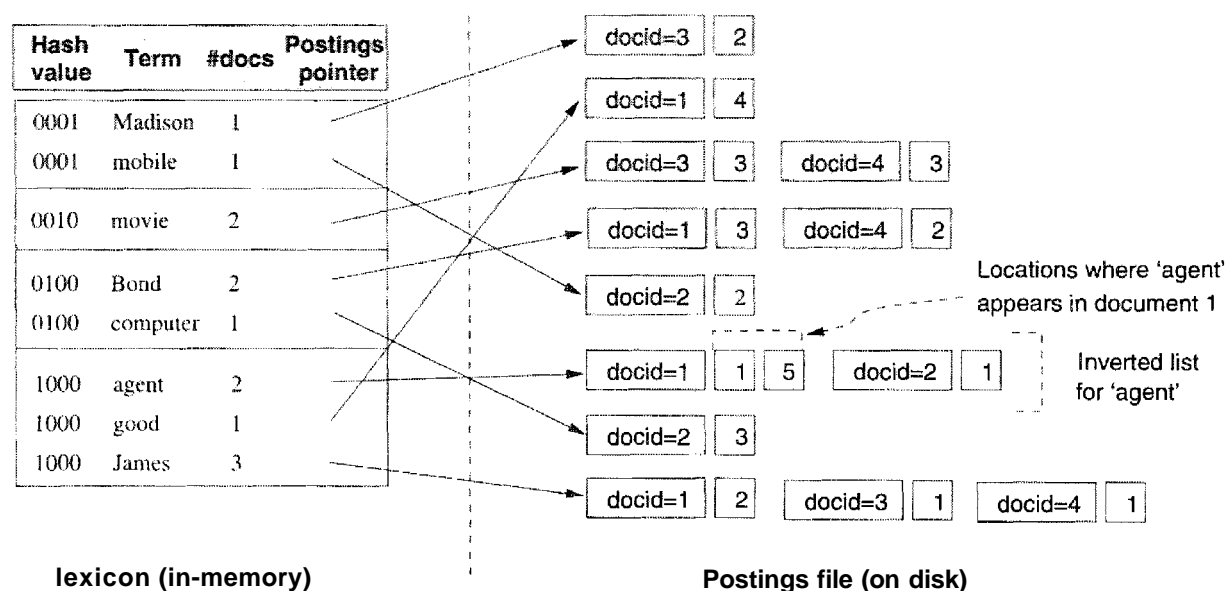


Figure 27.5 Inverted Index for Example Collection

list (i.e., the number of documents that the term appears in). In general, it could contain additional information such as the IDF for the term, but it is important to keep the entry's size as small as possible.

The lexicon is maintained in-memory, and enables fast retrieval of the inverted list for a query term. The lexicon in Figure 27.5 uses a hash index, and is sketched by showing the hash value for the term; entries for terms are grouped into hash buckets by their hash value.

Using an Inverted Index

A query containing a single term is evaluated by first searching the lexicon to find the address of the inverted list for the term. Then the inverted list is retrieved, the docids in it are mapped to physical document addresses, and the corresponding documents are retrieved. If the results are to be ranked, the relevance of each document in the inverted list to the query term is computed, and documents are then retrieved in order of their relevance rank. Observe that the information needed to compute the relevance measure described in Section 27.2—the frequency of the query term in the document, the IDF of the term in the document collection, and the length of the document if it is used for length normalization—are all available in either the lexicon or the inverted list.

When inverted lists are very long, as in Web search engines, it is useful to consider whether we should precompute the relevance of each document in the inverted list for a term (with respect to that term) and sort the list by relevance rather than document id. This would speed up querying because we can just

look at a prefix of the inverted list, since users rarely look at 111010 than the first few results. However, maintaining lists in sorted order by relevance can be expensive. (Sorting by document id is convenient because new documents are assigned increasing ids, and we can therefore simply append entries for new documents at the end of the inverted list. Further, if the similarity function is changed, we do not have to rebuild the index.)

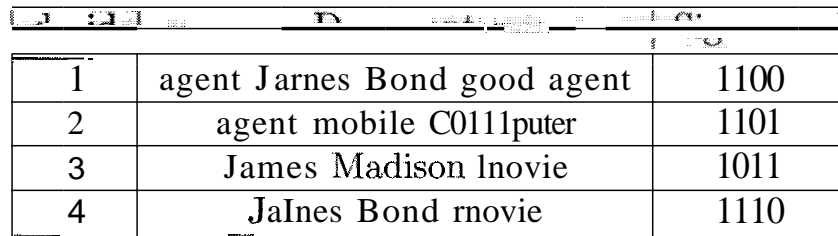
A query with a conjunction of several terms is evaluated by retrieving the inverted lists of the query terms one at a time and intersecting them. In order to minimize memory usage, the inverted lists should be retrieved in order of increasing length. A query with a disjunction of several terms is evaluated by merging all relevant inverted lists.

Consider the example inverted index shown in Figure 27.5. To evaluate the query 'James', we probe the lexicon to find the address of the inverted list for 'James', fetch it from disk and then retrieve document 1. To evaluate the query 'James' AND 'Bond', we first retrieve the inverted list for the term 'Bond' and intersect it with the inverted list for the term 'James.' (The inverted list of the term 'Bond' has length two, whereas the inverted list of the term 'James' has length three.) The result of the intersection of the list (1,4) with the list (1,3,4) is the list (1,4) and documents 1 and 4 are therefore retrieved. To evaluate the query 'James' OR 'Bond,' we retrieve the two inverted lists in any order and merge the results.

For ranked queries with multiple terms, we must fetch the inverted lists for all terms, compute the relevance of every document that appears in one of these lists with respect to the given collection of query terms, and then sort the document ids by their relevance before fetching the documents in relevance rank order. Again, if the inverted lists are sorted by the relevance measure, we can support ranked queries by typically processing only small prefixes of the inverted lists. (Observe that the relevance of a document with respect to the query is easily computed from its relevance with respect to each query term.)

27.3.2 Signature Files

A signature file is another index structure for text database systems that supports efficient evaluation of boolean queries. A signature file contains an index record for each document in the database. This index record is called the signature of the document. Each signature has a fixed size of b bits; b is called the signature width. The bits that are set depend on the words that appear in the document. We map words to bits by applying a hash function to each word in the document and we set the bits that appear in the result of



1	agent Jarnes Bond good agent	1100
2	agent mobile C0111puter	1101
3	James Madison Inovie	1011
4	JaInes Bond movie	1110

Figure 27.6 Signature File for Example Collection

the hash function. Note that unless we have a bit for each possible word in the vocabulary, the same bit could be set twice by different words because the hash function maps both words to the same bit. We say that a signature S_1 matches another signature S_2 if all the bits that are set in signature S_2 are also set in signature S_1 . If signature S_1 matches signature S_2 , then signature S_1 has at least as many bits set as signature S_2 .

For a query consisting of a conjunction of terms, we first generate the query signature by applying the hash function to each word in the query. We then scan the signature file and retrieve all documents whose signatures match the query signature, because every such document is a potential result to the query. Since the signature does not uniquely identify the words that a document contains, we have to retrieve each potential match and check whether the document actually contains the query terms. A document whose signature matches the query signature but that does not contain all terms in the query is called a false positive. A false positive is an expensive mistake since the document has to be retrieved from disk, parsed, stemmed, and checked to determine whether it contains the query terms.

For a query consisting of a disjunction of terms, we generate a list of query signatures, one for each term in the query. The query is evaluated by scanning the signature file to find documents whose signatures match any signature in the list of query signatures.

As an example, consider the signature file of width 4 for our running example shown in Figure 27.6. The bits set by the hashed values of all query terms are shown in the figure. To evaluate the query 'James,' we first compute the hash value of the term; this is 1000. Then we scan the signature file and find matching index records. As we can see from Figure 27.6, the signatures of all records have the first bit set. We retrieve all documents and check for false positives; the only false positive for this query is document with rid 2. (Unfortunately, the hashed value of the term 'agent' also happened to set the very first bit in the signature.) Consider the query 'James' And 'Bond.' The query signature is 1100 and three document signatures match the query signature. Again, we retrieve one false positive. As another example of a conjunctive query, con-

sider the query ‘movie’ And ‘Madison.’ The query signature is 0011, and only one document signature matches the query signature. No false positives are retrieved.

Note that for each query we have to scan the complete signature file, and there are as many records in the signature file as there are documents in the database. To reduce the amount of data that has to be retrieved for each query, we can vertically partition a signature file into a set of bit slices, and we call such an index a bit-sliced signature file. The length of each bit slice is still equal to the number of documents in the database, but for a query with q bits set in the query signature we need only to retrieve q bit slices. The reader is invited to construct a bit-sliced signature file and to evaluate the example queries in this paragraph using the bit slices.

27.4 WEB SEARCH ENGINES

Web search engines must contend with extremely large numbers of documents, and have to be highly scalable. Documents are also linked to each other, and this link information turns out to be very valuable in finding pages relevant to a given search. These factors have caused search engines to differ from traditional IR systems in important ways. Nonetheless, they rely on some form of inverted indexes as the basic indexing mechanism. In this section, we discuss Web search engines, using Google as a typical example.

27.4.1 Search Engine Architecture

Web search engines crawl the web to collect documents to index. The crawling algorithm is simple, but crawler software can be complex because of the details of connecting to millions of sites, minimizing network latencies, parallelizing the crawling, dealing with timeouts and other connection failures, ensuring that crawled sites are not unduly stressed by the crawler, and other practical concerns.

The search algorithm used by a crawler is a graph traversal. Starting at a collection of pages with many links (e.g., Yahoo directory pages), all links on crawled pages are followed to identify new pages. This step is iterated, keeping track of which pages have been visited in order to avoid re-visiting them.

The collection of pages retrieved through crawling can be enormous, on the order of billions of pages. Indexing them is a very expensive task. Fortunately, the task is highly parallelizable: Each document is independently analyzed to create inverted lists for the terms that appear in the document. These per-document lists are then sorted by term and merged to create complete per-

term inverted lists that span all documents. Term statistics such as IDF can be computed during the merge phase.

Supporting searches over such vast indexes is another nontrivial undertaking. Fortunately, again, the task is readily parallelized using a cluster of inexpensive machines: We can deal with the amount of data by partitioning the index across several machines. Each machine contains the inverted index for those terms that are mapped to that machine (e.g., by hashing the term). Queries may have to be sent to multiple machines if the terms they contain are handled by different machines, but given that Web queries rarely contain more than two terms, this is not a serious problem in practice.

We must also deal with a huge volume of queries; Google supports over 150 million searches each day, and the number is growing. This is accomplished by replicating the data across several machines. We already described how the data is partitioned across machines. For each partition, we now assign several machines, each of which contains an exact copy of the data for that partition. Queries on this partition can be handled by any machine in the partition. Queries can be distributed across machines on the basis of load, by hashing on IP addresses, etc. Replication also addresses the problem of high-availability, since the failure of a machine only increases the load on the remaining machines in the partition, and if partitions contain several machines the impact is small. Failures can be made transparent to users by routing queries to other machines through the load balancer.

27.4.2 Using Link Information

webpages are created by a variety of users for a variety of purposes, and their content does not always lend itself to effective retrieval. The most relevant pages for a search may not contain the search terms at all and are therefore not returned by a boolean keyword search! For example, consider the query term 'Web browser.' A boolean text query using the terms does not return the relevant pages of Netscape Corporation or Microsoft, because these pages do not contain the term 'Web browser' at all. Similarly, the home page of 'Yahoo' does not contain the term 'search engine.' The problem is that relevant sites do not necessarily describe their contents in a way that is useful for boolean text queries.

Until now, we only considered information within a single webpage to estimate its relevance to a query. But webpages are connected through hyperlinks, and it is quite likely that there is a webpage containing the term 'search engine' that has a link to Yahoo's home page. Can we use the information hidden in such links?

Building on research in the sociology literature, an interesting analogy between links and bibliographic citations suggests a way to exploit link information: Just as influential authors and publications are cited often, good webpages are likely to be often linked to. It is useful to distinguish between two types of pages, *authorities* and *hubs*. An authority is a page that is very relevant to a certain topic and that is recognized by other pages as authoritative on the subject. These other pages, called hubs, usually have a significant number of hyperlinks to authorities, although they themselves are not very well known and do not necessarily carry a lot of content relevant to the given query. Hub pages could be compilations of resources about a topic on a site for professionals, lists of recommended sites for the hobbies of an individual user, or even a part of the bookmarks of an individual user that are relevant to one of the user's interests; their main property is that they have many outgoing links to relevant pages. Good hub pages are often not well known and there may be few links pointing to a good hub. In contrast, good authorities are 'endorsed' by many good hubs and thus have many links from good hub pages.

This symbiotic relationship between hubs and authorities is the basis for the HITS algorithm, a link-based search algorithm that discovers high-quality pages that are relevant to a user's query terms. The HITS algorithm models Web as a directed graph. Each webpage represents a node in the graph, and a hyperlink from page *A* to page *B* is represented as an edge between the two corresponding nodes.

Assume that we are given a user query with several terms. The algorithm proceeds in two steps. In the first step, the *sampling step*, we collect a set of pages called the base set. The base set most likely includes very relevant pages to the user's query, but the base set can still be quite large. In the second step, the *iteration step*, we find good authorities and good hubs among the pages in the base set.

The sampling step retrieves a set of webpages that contain the query terms, using some traditional technique. For example, 'we can evaluate the query as a boolean keyword search and retrieve all webpages that contain the query terms. We call the resulting set of pages the root set. The root set might not contain all relevant pages because some authoritative pages might not include the user query words. But we expect that at least some of the pages in the root set contain hyperlinks to the most relevant authoritative pages or that some authoritative pages link to pages in the root set. This motivates our notion of a link page. We call a page a link page if it has a hyperlink to some page in the root set or if a page in the root set has a hyperlink to it. In order not to miss potentially relevant pages, we augment the root set by all link pages and we call the resulting set of pages the base set. Thus, the base set includes all

root pages and all link pages; we refer to a webpage in the base set as a base page.

Our goal in the second step of the algorithm is to find out which base pages are good hubs and good authorities and to return the best authorities and hubs as the answers to the query. To quantify the quality of a base page as a hub and as an authority, we associate with each base page in the base set a hub weight and an authority weight. The hub weight of the page indicates the quality of the page as a hub, and the authority weight of the page indicates the quality of the page as an authority. We compute the weights of each page according to the intuition that a page is a good authority if many good hubs have hyperlinks to it, and that a page is a good hub if it has many outgoing hyperlinks to good authorities. Since we do not have any a priori knowledge about which pages are good hubs and authorities, we initialize all weights to one. We then update the authority and hub weights of base pages iteratively as described below.

Consider a base page p with hub weight h_p and with authority weight a_p . In one iteration, we update a_p to be the sum of the hub weights of all pages that have a hyperlink to p . Formally:

$$a_p = \sum_{\text{All base pages } q \text{ that have a link to } p} h_q$$

Analogously, we update h_p to be the sum of the weights of all pages that p points to:

$$h_p = \sum_{\text{All base pages } q \text{ such that } p \text{ has a link to } q} a_q$$

Comparing the algorithm with the other approaches to querying text that we discussed in this chapter, we note that the iteration step of the HITS algorithm—the distribution of the weights—does not take into account the words on the base pages. In the iteration step, we are only concerned about the relationship between the base pages as represented by hyperlinks.

The HITS algorithm usually produces very good results. For example, the five highest ranked results from Google (which uses a variant of the HITS algorithm) for the query ‘Raghu Ramakrishnan’ are the following webpages:

```
www.cs.wisc.edu/~raghu/raghu.html
www.cs.wisc.edu/~dbbook/dbbook.html
www.informatik.uni-trier.de/
    ~ley/db/indices/a-tree/r/Ramakrishnan:Raghu.html
www.informatik.uni-trier.de/
```

Computing hub and authority weights: We can use matrix notation to write the updates for all hub and authority weights in one step. Assume that we have n web pages in the base set $\{1, 2, \dots, n\}$. The adjacency matrix B of the base set is an $n \times n$ matrix whose entries are either 0 or 1. The matrix entry (i, j) is set to 1 if page i has a hyperlink to page j ; it is set to 0 otherwise. We can also write the hub weights h and authority weights a in vector notation: $h = (h_1, \dots, h_n)$ and $a = (a_1, \dots, a_n)$. We can now rewrite our update rules as follows:

$$h = B \cdot a, \quad \text{and} \quad a = B^T \cdot h.$$

Unfolding this equation once, corresponding to the first iteration, we obtain:

$$h = BB^T h = (BB^T)h, \quad \text{and} \quad a = BTBa = (BTB)a.$$

After the second iteration, we arrive at:

$$h = (BB^T)^2 h, \quad \text{and} \quad a = (B^T B)^2 a.$$

Results from linear algebra tell us that the sequence of iterations for the hub (resp. authority) weights converges to the principal eigenvectors of BB^T (resp. $B^T B$) if we normalize the weights before each iteration so that the sum of the squares of all weights is always 2 \cdot n . Furthermore, results from linear algebra tell us that this convergence is independent of the choice of initial weights, as long as the initial weights are positive. Thus, our rather arbitrary choice of initial weights---we initialized all hub and authority weights to 1---does not change the outcome of the algorithm.

Google's Pigeon Rank: Google computes the *pigeon rank* (PR) for a webpage A using the following formula, which is very similar to the Hub-Authority ranking functions:

$$PR(A) = (1 - d) + d(PR(T_1)/C(T_1) + \dots + PR(T_n)/C(T_n))$$

$T_1 \dots T_n$ are the pages that link (or 'point') to A , $C(T_i)$ is the number of links going out of page T_i , and d is a heuristically chosen constant (Google uses 0.85). Pigeon ranks form a probability distribution over all webpages; the sum of ranks over all pages is 1. If we consider a model of user behavior in which a user randomly chooses a page and then repeatedly clicks on links until he gets bored and randomly chooses a new page, the probability that the user visits a page is its Pigeon rank. The pages in the result of a search are ranked using a combination of an IR-style relevance metric and Pigeon rank.

SQL/MM: Full Text 'Full text' is described as data that can be searched, unlike simple character strings, and a new data type called `FullText` is introduced to support it. The methods associated with this type support searching for individual words, phrases, words that 'sound like' a query term, etc. Three methods are of particular interest. `CONTAINS` checks if a `FullText` object contains a specified search term (word or phrase). `RANK` returns the relevance rank of a `FullText` object with respect to a specified search term. (How the rank is defined is left to the implementation.) `IS ABOUT` determines whether the `FullText` object is sufficiently related to the specified search term. (The behavior of `IS ABOUT` is also left to the implementation.)

Relational DBMSs from IBM, Microsoft, and Oracle all support text fields, although they do not currently conform to the SQL/MM standard.

[-ley/db/indices/a-tree/s/Seshadri:Praveen.html](http://ley/db/indices/a-tree/s/Seshadri:Praveen.html)

www.acm.org/awards/fellows_citations_n-z/ramakrishnan.html

The first result is Ramakrishnan's home page; the second is the home page for this book; the third is the page listing his publications in the popular DBLP bibliography; and the fourth (initially puzzling) result is the list of publications for a former student of his.

27.5 MANAGING TEXT IN A DBMS

In preceding sections, we saw how large text collections are indexed and queried in IR systems and Web search engines. We now consider the additional challenges raised by integrating text data into database systems.

The basic approach being pursued by the SQL standards community is to treat text documents as a new data type, `FullText`, that can appear as the value of a field in a table. If we define a table with a single column of type `FullText`, each row in the table corresponds to a document in a document collection. Methods of `FullText` can be used in the `WHERE` clause of SQL queries to retrieve rows containing text objects that match an IR-style search criterion. The relevance rank of a `FullText` object can be explicitly retrieved using the `RANK` method, and this can be used to sort results by relevance.

Several points must be kept in mind as we consider this approach:

- This is an extremely general approach, and the performance of a SQL system that supports such an extension is likely to be inferior to a specialized IR System.

- The model of data does not adequately reflect documents with additional metadata. If we store documents in a table with a FullText column and use additional columns to store metadata—for example, author, title, Summary, rating, popularity—relevance measures that combine metadata with IR similarity measures must be expressed using user-defined methods, because the RANK method only has access to the FullText object, and not the metadata. The emergence of XML documents, which have non-uniform, partial metadata, further complicates matters.
- The handling of updates is unclear. As we have seen, IR indexes are complex, and expensive to maintain. Requiring a system to update the indexes before the updating transaction commits can impose a severe performance penalty.

27.5.1 Loosely Coupled Inverted Index

The implementation approach used in current relational DBMSs that support text fields is to have a separate text-search engine that is loosely coupled to the DBMS. The engine periodically updates the indexes, but provides no transactional guarantees. Thus, a transaction could insert (a row containing) a text object and commit, and a subsequent transaction that issues a matching search might not retrieve the (row containing the) object.

27.6 A DATA MODEL FOR XML

As we saw in Section 7.4.1, XML provides a way to mark up a document with meaningful tags that impart some partial structure to the document. *Semistructured data models*, which we introduce in this section, capture much of the structure in XML documents, while abstracting away many details.¹ Semistructured data models have the potential to serve as a formal foundation for XQL and enable us to rigorously define the semantics of queries over XQL, which we discuss in Section 27.7.

27.6.1 Motivation for Loose Structure

Consider a set of documents on the Web that contain hyperlinks to other documents. These documents, although not completely unstructured, cannot be modeled naturally in the relational data model because the pattern of hyperlinks is not regular across documents. In fact, every HTML document has

¹An important aspect of XML that is *not* captured is the ordering of elements. A more complete data model called XData has been proposed by the W3C committee that is developing XML standards, but we do not discuss it here.

XML Data Models: A number of data models for XML are being considered by standards committees such as ISO and W3C. W3C's Infoset is a tree-structured model, and each node can be retrieved through an accessor function. A version called Post-Validation Infoset (PSVI) serves as the data model for XML Schema. The XQuery language has yet another data model associated with it. The plethora of models is due to parallel development in some cases, and due to different objectives in others. Nonetheless, all these models have loosely-structured trees as their central feature.

some minimal structure, such as the text in the TITLE tag versus the text in the document body, or text that is highlighted versus text that is not. As another example, a bibliography file also has a certain degree of structure due to fields such as *author* and *title*, but is otherwise unstructured text. Even data that is 'unstructured', such as free text or an image or a video clip, typically has some associated information such as timestamp or author information that contributes partial structure.

We refer to data with such partial structure as semistructured data. There are many reasons why data might be semistructured. First, the structure of data might be implicit, hidden, unknown, or the user might choose to ignore it. Second, when integrating data from several heterogeneous sources, data exchange and transformation are important problems. We need a highly flexible data model to integrate data from all types of data sources including flat files and legacy systems; a structured data model such as the relational model is often too rigid. Third, we cannot query a structured database without knowing the schema, but sometimes we want to query the data without full knowledge of the schema. For example, we cannot express the query "Where in the database can we find the string *Malgudi*?" in a relational database system without knowing the schema, and knowing which fields contain such text values.

27.6.2 A Graph Model

All data models proposed for semistructured data represent the data as some kind of labeled graph. Nodes in the graph correspond to compound objects or atomic values. Each edge indicates an object-subobject or object-value relationship. Leaf nodes, i.e., nodes with no outgoing edges have a value associated with them. There is no separate schema and no auxiliary description; the data in the graph is self-describing. For example, consider the graph shown in Figure 27.7, which represents part of the XML data from Figure 7.2. The root node of the graph represents the outermost element, BOOKLIST. The node has three children that are labeled with the element name BOOK, since the list of books

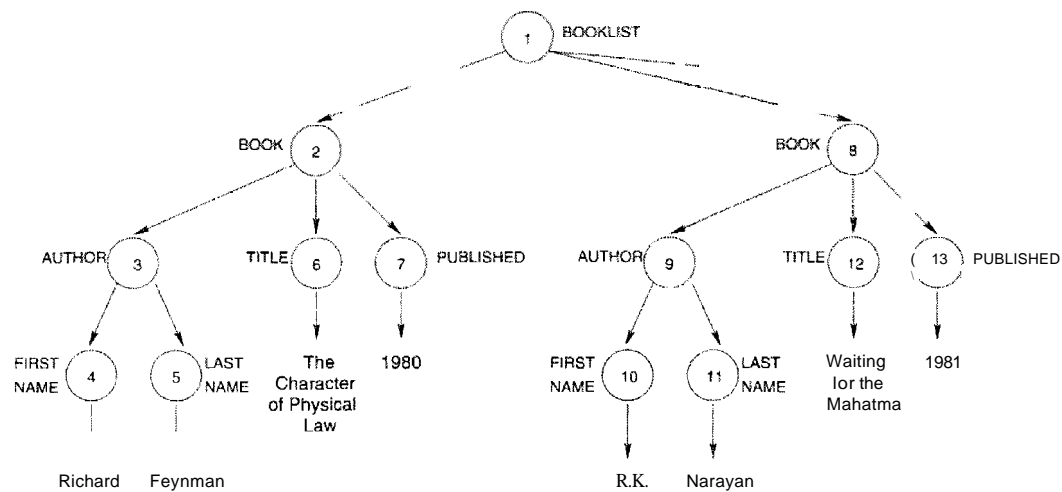


Figure 27.7 The Semistructured Data Model

consists of three individual books. The numbers within the nodes indicate the object identifier associated with the corresponding object.

We now describe one of the proposed data models for semistructured data, called the object exchange model (OEM). Each object is described by a quadruple consisting of a *label*, a *type*, the *value* of the object, and all *object identifier* which is a unique identifier for the object. Since each object has a label that can be thought of as a column name in the relational model, and each object has a type that can be thought of as the column type in the relational model, the object exchange model is self-describing. Labels in OEM should be as informative as possible, since they serve two purposes—they can be used to identify an object as well as to convey the meaning of an object. For example, we can represent the last Name of an author as follows:

(lastName, string, "Feynman")

More complex objects are decomposed hierarchically into smaller objects. For example, an author name can contain a first name and a last name. This object is described as follows:

(authorName, set, {firstName₁, lastName₁})
 firstName₁ is (firstName, string, "Richard")
 lastName₁ is (lastName, string, "Feynman")

As another example, an object representing a set of books is described as follows:

(bookList, set, {book₁, book₂, book₃})
 book_i is (book, set, {author_i, title_i, published_i})

SQL and XML: XQuery is a standard proposed by the World-Wide Web Consortium (W3C). In parallel, standards committees developing the SQL standards have been working on a successor to SQL:1999 that supports XML. The part that relates to XML is tentatively called SQL/XML and details can be found at <http://sqlx.org>.

```
book2 is ⟨book, set, {author2, title2, published2}⟩
book3 is ⟨book, set, {author3, title3, Published3}⟩
  author3 is (author, set, {first3, lastname3} )
  title3 is (title, string, liThe English Teacher")
  published3 is (published, integer, 1980)
```

27.7 XQUERY: QUERYING XML DATA

Given that XML documents are encoded in a way that reflects (a considerable amount of) structure, we have the opportunity to use a high-level language that exploits this structure to conveniently retrieve data from within such documents. Such a language would also allow us to easily translate XML data between different DTDs, as we illustrate when integrating data from multiple sources. At the time of writing of this book, XQuery is the W3C standard query language for XML data. In this section, we give a brief overview of XQuery.

27.7.1 Path Expressions

Consider the XML document shown in Figure 7.2. The following example query returns the last names of all authors, assuming that our XML document resides at the location `www.ourbookstore.com/books.xml`.

```
FOR
  $1 IN doc(www.ourbookstore.com/books.xml)//AUTHOR/LASTNAME
RETURN <RESULT> $1 </RESULT>
```

This example illustrates some of the basic constructs of XQuery. The FOR clause in XQuery is roughly analogous to the FROM clause in SQL. The RETURN clause is similar to the SELECT clause. We return to the general form of queries shortly, after introducing an important concept called a path expression.

The expression

```
doc(www.ourbookstore.com/books.xml)//AUTHOR/LASTNAME
```

XPath and Other XML Query Languages: Path expressions in XQuery are derived from XPath, an earlier XML query facility. Path expressions in XPath can be qualified with selection conditions, and can utilize several built-in functions (e.g., counting the number of nodes matched by the expression). Many of XQuery's features are borrowed from earlier languages, including XML-QL and Quilt.

in the FOR clause is an example of a path expression. It specifies a path involving three entities: the document itself, the AUTHOR elements and the LASTNAME elements.

The path relationship is expressed through separators / and //. The separator // specifies that the AUTHOR element can be nested anywhere within the document whereas the separator / constrains the LASTNAME element to be nested immediately under (in terms of the graph structure of the document) the AUTHOR element. Evaluating a path expression returns a *set* of elements that match the expression. The variable *i* in the example query is bound in turn to each LASTNAME element returned by evaluating the path expression. (To distinguish variable names from normal text, variable names in XQuery are prefixed with a dollar sign \$.)

The RETURN clause constructs the query result—which is also an XML document—by bracketing each value to which the variable *i* is bound with the tag RESULT. If the example query is applied to the sample data shown in Figure 7.2, the result would be the following XML document:

```
<RESULT><LASTNAME>Feynman </LASTNAME></RESULT>
<RESULT><LASTNAME>Narayan </LASTNAME></RESULT>
```

We use the document in Figure 7.2 as our input in the rest of this chapter.

27.7.2 FLWR Expressions

The basic form of an XQuery consists of a **FLWR expression**, where the letters denote the FOR, LET, WHERE and RETURN clauses. The FOR and LET clauses bind variables to values through path expressions. These values are qualified by the WHERE clause, and the result XML fragment is constructed by the RETURN clause.

The difference between a FOR and LET clause is that while FOR binds a variable to each element specified by the path expression, LET binds a variable to the whole *collection* of elements. Thus, if we change our example query to:


```
LET
  $IINdoc(www.ourbookstore.com/books.xml)//AUTHOR/LASTNAME
RETURN <RESULT> $1 </RESULT>
```

then the result of the query beconles:

```
<RESULT>
  <LASTNAME>Feynman</LASTNAME>
  <LASTNAME>Narayan</LASTNAME>
</RESULT>
```

Selection conditions are expressed using the WHERE clause. Also, the output of a query is not lirnited to a single elernent. These points are illustrated by the following query, which finds the first and last names of all authors who wrote a book that was published in 1980:

```
FOR $b IN doc(www.ourbookstore.com/books.xml)/BOOKLIST/BOOK
WHERE $b/PUBLISHED='19S0'
RETURN
  <RESULT> $b/AUTHOR/FIRSTNAME, $b/AUTHOR/LASTNAME </RESULT>
```

The result of the above query is the following XML docurnent:

```
<RESULT>
  <FIRSTNAME>Richard </FIRSTNAME><LASTNAME>Feynman </LASTNAME>
</RESULT>
<RESULT>
  <FIRSTNAME>R.K. </FIRSTNAME><LASTNAME>Narayan </LASTNAME>
</RESULT>
```

For the specific DTI) in this exalnple, where a BOOK elernent **has** only one AUTHOR, the above query can be written by using a different path expression in the FOR clause, as follows.

```
FOR $a IN
  doc(www.ourbookstore.com/books.xml)
  /BOOKLIST/BOOK[PUBLISHED='19S0']/AUTHOR
RETURN <RESULT> $a/FIRSTNAME, $a/LASTNAME </RESULT>
```

rrhe path expression in this query is an instance of a **branching path expression**. The variable *l* is now bound to every AUTHOR elernent that rnatches the path doc/BOOKLIST/BOOK/AUTHOR where the intennediate BOOK elClnent is constrained to have a PUBLISHED elernent nested inunediatly within it with the value 1980.

27.7.3 Ordering of Elements

XML data consists of *ordered* documents and so the query language XQuery returns data in source order. The semantics of XQuery is that a path expression returns results sorted in document order. Thus, variables in the FOR clause are bound in document order. If however, we desire a different order, we can explicitly order the output as shown in the following query, which returns TITLE elements sorted lexicographically.

```
FOR
  $b IN doc(www.ourbookstore.com/books.xml)/BOOKLIST/BOOK
RETURN <BOOKTITLES> $b/TITLE </BOOKTITLES>
SORT BY TITLE
```

27.7.4 Grouping and Generation of Collection Values

Our next example illustrates grouping in XQuery, which allows us to generate a new collection value for each group. (Contrast this with grouping in SQL, which only allows us to generate an aggregate value (e.g., SUM) per group.) Suppose that for each year we want to find the last names of authors who wrote a book published in that year. We group by year of publication and generate a list of last names for each year:

```
FOR $p IN DISTINCT
  doc(www.ourbookstore.com/books.xml)/BOOKLIST/BOOK/PUBLISHED
RETURN
  <RESULT>
    $p,
    FOR $a IN DISTINCT /BOOKLIST/BOOK[PUBLISHED=$p]/AUTHOR
    RETURN $a
  </RESULT>
```

The keyword *DISTINCT* eliminates duplicates from the collection returned by a path expression. Using the XML document in Figure 7.2 as input, the above query produces the following result:

```
<RESULT> <PUBLISHED>1980</PUBLISHED>
  <LASTNAME>Feynman</LASTNAME>
  <LASTNAME>Narayan</LASTNAME>
</RESULT>
<RESULT> <PUBLISHED>1981</PUBLISHED>
  <LASTNAME>Narayan</LASTNAME>
</RESULT>
```

27.8 EFFICIENT EVALUATION OF XML QUERIES

XQuery operates on XML data and produces XTvL data as output. In order to be able to evaluate queries efficiently, we need to address the following issues.

- **Storage:** We can use an existing storage system like a relational or object oriented system or design a new storage format for XML documents. There are several ways to use a relational system to store XML. One of them is to store the XML data as Character Large Objects (CLOBs). (CLOBs were discussed in Chapter 23.) In this case, however, we cannot exploit the query processing infrastructure provided by the relational system and would instead have to process XQuery outside the database system. In order to circumvent this problem, we need to identify a schema according to which the XML data can be stored. These points are discussed in Section 27.8.1.
- **Indexing:** Path expressions add a lot of richness to XQuery and yield many new access patterns over the data. If we use a relational system for storing XML data, then we are constrained to use only relational indexes like the B-Tree. However, if we use a native storage engine, then we have the option of building novel index structures for path expressions, some of which are discussed in Section 27.8.2.
- **Query Optimization:** Optimization of queries in XQuery is an open problem. The work so far in this area can be divided into three parts. The first is developing an algebra for XQuery, analogous to relational algebra. The second research direction is providing statistics for path expression queries. Finally, some work has addressed simplification of queries by exploiting constraints on the data. Since query optimization for XQuery is still at a preliminary stage, we do not cover it in this chapter.

Another issue to be considered while designing a new storage system for XML data is the verbosity of repeated tags. As we see in Section 27.8.1) using a relational storage system addresses this problem since tag names are not stored repeatedly. If on the other hand, we want to build a native storage system, then the manner in which the XML data is compressed becomes significant. Several compression algorithms are known that achieve compression ratios close to relational storage, but we do not discuss them here.

27.8.1 Storing XML in RDBMS

One natural candidate for storing XML data is a relational database system. The main issues involved in storing XML data in a relational system are:

Commercial database systems and XML: Many relational and object-relational database system vendors are currently looking into support for XML in their database engines. Several vendors of object-oriented database management systems already offer database engines that can store XML data whose contents can be accessed through graphical user interfaces or server-side Java extensions.

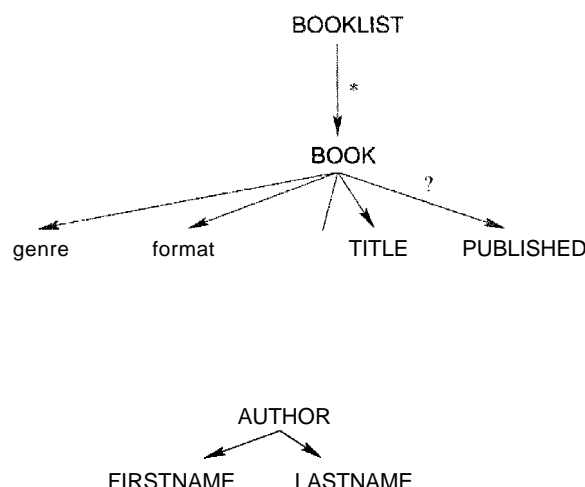


Figure 27.8 Bookstore XML DTD Element Relationships

- ii *Choice of relational schema:* In order to use an RDBMS, we need a schema. What relational schema should we use even assuming that the XML data comes with an associated schema?
- ii *Queries:* Queries on XML data are in XQuery whereas a relational system can only handle SQL. Queries in XQuery therefore need to be *translated* into SQL.
- *Reconstruction:* The output of XQuery is XML. Thus, the result of a SQL query needs to be converted back into XML.

Mapping XML Data to Relations

We illustrate the mapping process through our bookstore example. The nesting relationships among the different elements in the DTD is shown in Figure 27.8. The edges indicate the nature of the nesting.

One way to derive a relational schema is as follows. We begin at the BOOKLIST element and create a relation to store it. Traversing down from BOOKLIST, we get BOOK following a *** edge. This edge indicates that we store the BOOK elements in a separate relation. Traversing further down, we see that all elements and

attributes nested within BOOK occur at most once. Hence, we can store them in the same relation as BOOK. The resulting relational schema *RelSchema1* is shown below.

```
BOOKLIST(id: integer)
BOOK(booklistid: integer, author_firstname: string,
     author_lastname: string, title: string,
     published: string, genre: string, format: string)
```

BOOK. *booklistid* connects BOOK to BOOKLIST. Since a DTD has only one base type, **string**, the only base type used in the above schema is **string**. The constraints expressed through the DTD are expressed in the relational schema. For instance, since every BOOK must have a TITLE child, we must constrain the *title* column to be *non-null*.

Alternatively, if the DTD is changed to allow BOOK to have more than one AUTHOR child, then the AUTHOR elements cannot be stored in the same relation as BOOK. This change yields the following relational schema *RelSchema2*.

```
BOOKLIST(id: integer)
BOOK(id: integer, booklistid: integer,
     title: string, published: string, genre: string, format: string)
AUTHOR(bookid: integer, firstname: string, lastname: string)
```

The column AUTHOR. *bookid* connects AUTHOR to BOOK.

Query Processing

Consider the following example query again:

```
FOR
  $b IN doc(www.ourbookstore.com/books.xml)/BOOKLIST/Book
WHERE $b/PUBLISHED='1980'
RETURN
  <RESULT> $b/AUTHOR/FIRSTNAME, $b/AUTHOR/LASTNAME </RESULT>
```

If the mapping between the XML data and relational tables is known, then we can construct a SQL query that returns all columns that are needed to reconstruct the result XML document for this query. Conditions enforced by the path expressions and the WHERE clause are translated into equivalent conditions in the SQL query. We obtain the following equivalent SQL query if we use *RelSchema1* as our relational schema.

```
SELECT BOOK.author_firstname, BOOK.author_lastname
```

```
FROM    BOOK, BOOKLIST
WHERE   BOOKLIST.id = BOOK.booklistid
        AND BOOK.published='1980'
```

The results thus returned by the relational query processor are then tagged, outside the relational system, as specified by the RETURN clause. This is the result of the *reconstruct'ion* phase.

In order to understand this better, consider what happens if we allow a BOOK to have multiple AUTHOR children. Assume that we use *Rel8chema2* as our relational schema. Processing the FOR and WHERE clauses tells us that it is necessary to join relations BOOKLIST and BOOK with a selection on the BOOK relation corresponding to the year condition in the above query. Since the RETURN clause needs information about AUTHOR elements, we need to further join the BOOK relation with the AUTHOR relation and project the *first'name* and *last'name* columns in the latter. Finally, since each binding of the variable \$b in the above query produces one RESULT element, and since each BOOK is now allowed to have more than one AUTHOR, we need to project the *id* column of the BOOK relation. Based on these observations, we obtain the following equivalent SQL query:

```
SELECT   BOOK.id, AUTHOR.firstname , AUTHOR.lastname
FROM     BOOK, BOOKLIST, AUTHOR
WHERE    BOOKLIST.id = BOOK.booklistid AND
        BOOK.id = AUTHOR.bookid AND BOOK.published='1980'
GROUP BY BOOK.id
```

The result is grouped by BOOK.id. The tagger outside the database system now receives results clustered by the BOOK element and can tag the resulting tuples on the fly.

Publishing Relational Data as XML

Since XML has emerged as the standard data exchange format for business applications, it is necessary to publish existing business data as XML. Most operational business data is stored in relational systems. Consequently, mechanisms have been proposed to publish such data as XML documents. These involve a language for specifying how to tag and structure relational data and an implementation to carry out the conversion. This mapping is in some sense the reverse of the XML-to-relational mapping used to store XML data. The conversion process mirrors the reconstruction phase when we execute XQuery using a relational system. The published XML data can be thought of as an XML view of relational data. This view can be queried using XQuery. One

One method of executing XQuery on such views is to translate the XML into SQL and then construct the XML result.

27.8.2 Indexing XML Repositories

Path expressions are at the heart of all proposed XML query languages, in particular XQuery. A natural question that arises is how to index XML data to support path expression evaluation. The aim of this section is to give a flavor of the indexing techniques proposed for this problem. We consider the OEM model of semi-structured data, where the data is self-describing and there is no separate schema.

Using a B+ Tree to Index Values

Consider the following XQuery example, which we discussed earlier on the bookstore XML data in Figure 7.2. The OEM representation of this data is shown in Figure 27.7.

```
FOR
  $b IN doc(www.ourbookstore.com/books.xml)/BOOKLIST/BOOK
WHERE $b/PUBLISHED='1980'
RETURN
  <RESULT> $b/AUTHOR/FIRSTNAME, $b/AUTHOR/LASTNAME </RESULT>
```

This query specifies joins along the objects with labels BOOKLIST, BOOK, AUTHOR, FIRSTNAME, LASTNAME and PUBLISHED with a selection condition on PUBLISHED objects.

Let us suppose that we are evaluating this query in the absence of any indexes for path expressions. However, we do have a value index such as a B-Tree that enables us to find the ids of all objects with label PUBLISHED and value 1980. There are several ways of executing this query under these assumptions.

For instance, we could begin at the document root and traverse down the data graph through the BOOKLIST object to the BOOK objects. By further traversing the data graph downwards, for each BOOK object we can check whether it satisfies the value predicate (PUBLISHED='1980'). Finally, for those BOOK objects that satisfy the predicate, we can find the relevant FIRSTNAME and LASTNAME objects. This approach corresponds to a top-down evaluation of the query.

Alternatively, we could begin by using the value index to find all PUBLISHED objects that satisfy PUBLISHED='1980'. If the data graph can be traversed in the reverse direction—that is, given an object, we can find its parent—then we

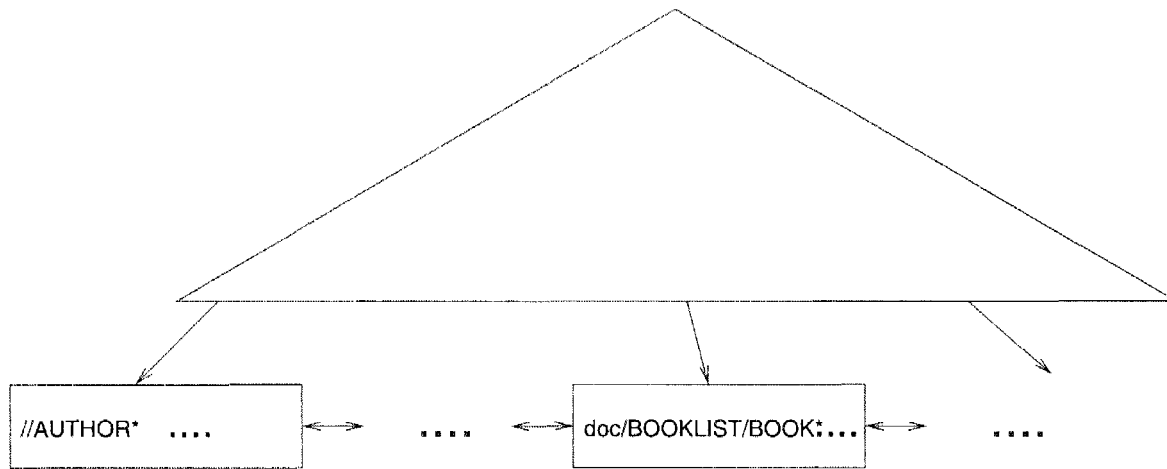


Figure 27.9 Path Expressions in a B-Tree

can find all parents of the PUBLISHED objects retaining only those that have label BOOK. We can continue in this manner until we find the FIRSTNAME and LASTNAME objects of interest. Observe that we need to perform all joins in the query on the fly.

Indexing on Structure vs. Value

Now let us ask ourselves whether traditional indexing solutions like the B-Tree can be used to index path expressions. We can use the B-Tree to map a path expression to the ids of all objects returned by it. The idea is to treat all path expressions as strings and order them lexicographically. Every leaf entry in the B-Tree contains a string representing a path expression and a list of ids corresponding to its result. Figure 27.9 shows how such a B-Tree would look. Let us contrast this with the traditional problem of indexing a well-ordered domain like integers for point queries. In the latter case, the number of distinct point queries that can be posed is just the number of data values and so is linear in the data size.

The scenario with path indexing is fundamentally different—the variety of ways in which we can combine tags to form (simple) path expressions coupled with the power of placing `//` separators leads to a much larger number of possible path expressions. For instance, an AUTHOR element in the example in Figure 27.7 is returned as part of the queries `BOOKLIST/BOOK/AUTHOR`, `//AUTHOR`, `//BOOK//AUTHOR`, `BOOKLIST//AUTHOR` and so on. The number of distinct queries can in fact be exponential in the data size (measured in terms of the number of XML elements) in the worst case. This is what motivates the search for alternative strategies to index path expressions.

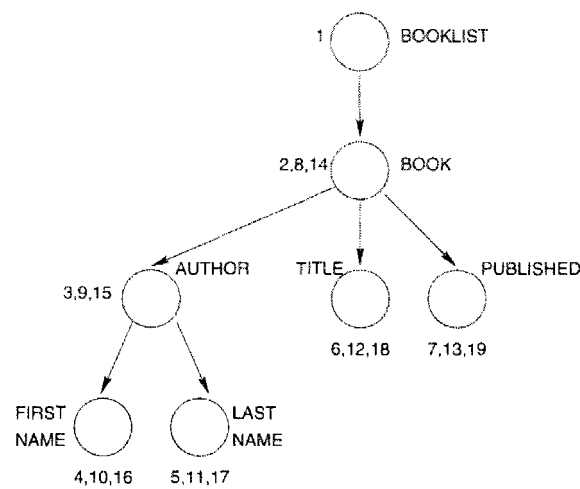


Figure 27.10 Example Path Index

The approach taken is to represent the mapping between a path expression and its result by means of a structural summary which takes the form of another labeled, directed graph. The idea is to preserve all the paths in the data graph in the summary graph, while having far fewer nodes and edges. An extent is associated with each node in the summary. The extent of an index node is a subset of the data nodes. The summary graph along with the extents constitutes a path index. A path expression is evaluated using the index by evaluating it against the summary graph and then taking the union of the extents of all matching nodes. This yields the index result of the path expression query. The index covers a path expression if the index result is the correct result; obviously, we can use an index to evaluate a path expression only if the index covers it.

Consider the structural summary shown in Figure 27.10. This is a path index for the data in Figure 27.7. The numbers shown beside the nodes correspond to the respective extents. Let us now examine how this index can change the top-down evaluation of the example query used earlier to illustrate B+ tree value indexes.

The top-down evaluation as outlined above begins at the document root and traverses down to the BOOK objects. This can be achieved more efficiently by the path index. Instead of traversing the data graph, we can traverse the path index down to the BOOK object in the index and look up its extent, which gives us the ids of all BOOK objects that match the path expression in the FOR clause. The rest of the evaluation then proceeds as before. Thus, the path index saves us from performing joins by essentially precomputing them. We note here that the path index shown in Figure 27.10 is isomorphic to the DTD schema graph shown in Figure 27.8. This drives home the point that the path index without the extents is a structural summary of the data.

the above path index is the Strong Dataguide. If we treat path expressions as strings, then the dataguide is the trie representing them. The trie is a well-known data structure used to search regular expressions over text. This shows the deeper unity between the research on indexing text and the XML path indexing work. Several other path indexes have been also proposed for semi-structured data, and this is an active area of research.

27.9 REVIEW QUESTIONS

Answers to the review questions can be found in the listed sections.

- What is information retrieval? (Section 27.1)
- What are some of the differences between DBMS and IR systems? Describe the differences between a ranked query and a boolean query. (Section 27.2)
- What is the vector space model, and what are its advantages? (Section 27.2.1)
- What is TF/IDF term weighting, and why do we weigh by both? We do we eliminate stop words? What is length normalization, and why is it done? (Section 27.2.2)
- How can we measure document similarity? (Sections 27.2.3)
- What are *precision* and *recall*, and how do they relate to each other? (Section 27.2.4)
- Describe the following two index structures for text: Inverted index and signature file. What is a bit-sliced signature file? (Section 27.3)
- How are web search engines architected? How does the “hubs and authorities” algorithm work? Can you illustrate it on a small set of pages? (Section 27.4)
- What support is there for managing text in a DBMS? (Section 27.5)
- Describe the OEM data model for semi-structured data. (Section 27.6)
- What are the elements of XQuery? What is a path expression? What is an FLWR expression? How can we order the output of query? How do we group query outputs? (Section 27.7)
- Describe how XML data can be stored in a relational DBMS. How do we map XML data to relations? Can we use the query processing infrastructure of the relational DBMS? How do we publish relational data as XML? (Section 27.8.1)

- How do we index collections of XML documents? What is the difference between indexing on structure versus indexing on value? What is a path index? (Section 27.8.2)

EXERCISES

Exercise 27.1 Carry out the following tasks.

1. Given an ASCII file, compute the frequency of each word and create a plot similar to Figure 27.3. (Feel free to use public domain plotting software.) Run the program on the collection of files currently in your directory and see whether the distribution of frequencies is Zipfian. How can you use such plots to create lists of stop words?
2. The Porter stemmer is widely used, and code implementing it is freely available. Download a copy, and run it on your collection of documents.
3. One criticism of the vector space model and its use in similarity checking is that it treats terms as occurring independently of each other. In practice, many words tend to occur together (e.g., ambulance and emergency). Write a program that scans an ASCII file and lists all pairs of words that occur within 5 words of each other. For each pair of words, you now have a frequency, and should be able to create a plot like Figure 27.3 with pairs of words on the X-axis. Run this program on some sample document collections. What do the results suggest about co-occurrences of words?

Exercise 27.2 Assume you are given a document database that contains six documents. After stemming, the documents contain the following terms:

Document	Terms
1	car manufacturer Honda auto
2	auto computer navigation
3	Honda navigation
4	manufacturer computer IBM
5	IBM personal computer
6	car Beetle VW

Answer the following questions.

1. Show the result of creating an inverted file on the documents.
2. Show the result of creating a signature file with a width of 5 bits. Construct your own hashing function that maps terms to bit positions.
3. Evaluate the following boolean queries using the inverted file and the signature file that you created: 'car', 'IBM' AND 'computer', 'IBM' AND 'car', 'IBM' OR 'auto', and 'IBM' AND 'computer' AND 'manufacturer'.
4. Assume that the query load against the document database consists of exactly the queries that were stated in the previous question. Also assume that each of these queries is evaluated exactly once.
 - (a) Design a signature file with a width of 3 bits and design a hashing function that minimizes the overall number of false positives retrieved when evaluating the

- (b) Design a signature file with a width of 6 bits and a hashing function that minimizes the overall number of false positives.
 - (c) Assume you want to construct a signature file. What is the smallest signature width that allows you to evaluate all queries without retrieving any false positives?
5. Consider the following ranked queries: 'car', 'IBM Computer', 'IBM car', 'IBM auto', and 'IBM Computer manufacturer'.
- (a) Calculate the IDF for every term in the database.
 - (b) For each document, show its document vector.
 - (c) For each query, calculate the relevance of each document in the database, with and without the length normalization step.
 - (d) Describe how you would use the inverted index to identify the top two documents that match each query.
 - (e) How would having the inverted lists sorted by relevance instead of document id affect your answer to the previous question?
 - (f) Replace each document with a variation that contains 10 copies of the same document. For each query, recompute the relevance of each document, with and without the length normalization step.

Exercise 27.3 Assume you are given the following structured document database:

Document	Terms
1	car car Manufacturer car car Honda auto
2	auto computer navigation
3	Honda navigation auto
4	manufacturer computer IBM/II graphics
5	IBM personal IBM computer IBM/II IBM/I! IBM IBM
6	car Beetle VW Honda

Using this database, repeat the previous exercise.

Exercise 27.4 You are in charge of the Genghis ('We execute fast') search engine. You are designing your server cluster to handle 500 million hits a day and 10 billion pages of indexed data. Each machine costs \$1000, and can store 10 million pages and respond to 200 queries per second (against these pages).

1. If you were given a budget of \$500,000 dollars for purchasing machines, and were required to index all 10 billion pages, could you do it?
2. What is the minimum budget to index all pages? If you assume that each query can be answered by looking at data in just one (10 million page) partition, and that queries are uniformly distributed across partitions, what peak load (in number of queries per second) can such a cluster handle?
3. How would your answer to the previous question change if each query, on average, accessed two partitions?
4. What is the minimum budget required to handle the desired load of 500 million hits per day if all queries are on a *single partition*? Assume that queries are uniformly distributed with respect to time of day.

5. How would your answer to the previous question change if the number of queries per day went up to 5 billion hits per day? How would it change if the number of pages went up to 100 billion?
6. Assume that each query accesses just one partition, that queries are uniformly distributed across partitions, but that at any given time, the peak load on a partition is up to 10 times the average load. What is the minimum budget for purchasing machines in this scenario?
7. Take the cost for machines from the previous question and multiply it by 10 to reflect the costs of maintenance, administration, network bandwidth, etc. This amount is your annual cost of operation. Assume that you charge advertisers 2 cents per page. What fraction of your inventory (i.e., the total number of pages that you serve over the course of a year) do you have to sell in order to make a profit?

Exercise 27.5 Assume that the base set of the HITS algorithm consists of the set of Web pages displayed in the following table. An entry should be interpreted as follows: Web page 1 has hyperlinks to pages 5 and 6.

Webpage	Pages that this page has links to
1	5, 6, 7
2	5, 7
3	6, 8
4	
5	1, 2
6	1, 3
7	1, 2
8	4

1. Run five iterations of the HITS algorithm and find the highest ranked authority and the highest ranked hub.
2. Compute Google's Pigeon Rank for each page.

Exercise 27.6 Consider the following description of items shown in the Eggface computer mail-order catalog.

"Eggface sells hardware and software. We sell the new Palm Pilot V for \$400; its part number is 345. We also sell the IBM ThinkPad 570 for only \$1999; its part number is 3784. We sell both business and entertainment software. Microsoft Office 2000 has just arrived and you can purchase the Standard Edition for only \$140, part number 974; the Professional Edition is \$200, part 975. The new desktop publishing software from Adobe called InDesign is here for only \$200, part 664. We carry the newest games from Blizzard software. You can start playing Diablo II for only \$30, part number 12, and you can purchase Starcraft for only \$10, part number 812. Our goal is complete customer satisfaction.....if we don't have what you want in stock, we'll give you \$10 off your next purchase!"

1. Design an HTML document that depicts the items offered by Eggface.
2. Create a well-formed XML document that describes the contents of the Eggface catalog.
3. Create a TTD for your XML document and make sure that the document you created in the last question is valid with respect to this TTD,

4. Write an XQuery query that lists all software items in the catalog, sorted by price.
5. Write an XQuery query that, for each vendor, lists all software items from that vendor (i.e., one row in the result per vendor).
6. Write an XQuery query that lists the prices of all hardware items in the catalog.
7. Depict the catalog data in the semistructured data model as shown in Figure 27.7.
8. Build a dataguide for this data. Discuss how it can be used (or not) for each of the above queries.
9. Design a relational schema to publish this data.

Exercise 27.7 A university database contains information about professors and the courses they teach. The university has decided to publish this information on the Web and you are in charge of the execution. You are given the following information about the contents of the database:

In the fall semester 1999, the course 'Introduction to Database Management Systems' was taught by Professor Ioannidis. The course took place Mondays and Wednesdays from 9–10 a.m. in room 101. The discussion section was held on Fridays from 9–10 a.m. Also in the fall semester 1999, the course 'Advanced Database Management Systems' was taught by Professor Carey. Thirty five students took that course which was held in room 110 Tuesdays and Thursdays from 1–2 p.m. In the spring semester 1999, the course 'Introduction to Database Management Systems' was taught by U.N. Owen on Tuesdays and Thursdays from 3–4 p.m. in room 110. Sixty three students were enrolled; the discussion section was on Thursdays from 4–5 p.m. The other course taught in the spring semester was 'Advanced Database Management Systems' by Professor Ioannidis, Monday, Wednesday, and Friday from 8–9 a.m.

1. Create a well-formed XML document that contains the university database.
2. Create a DTD for your XML document. Make sure that the XML document is valid with respect to this DTD.
3. Write an XQuery query that lists the names of all professors in the order they are listed on the Web.
4. Write an XQuery query that lists all courses taught in 1999. The result should be grouped by professor, with one row per professor, sorted by last name. For a given professor, courses should be ordered by title and should not contain duplicates (i.e., even if a professor teaches the same course twice in 1999, it should appear only once in the result).
5. Build a dataguide for this data. Discuss how it can be used (or not) for each of the above queries.
6. Design a relational schema to publish this data.
7. Describe the information in a different XML document—a document that has a different structure. Create a corresponding DTD and make sure that the document is valid. Re-formulate the queries you wrote for preceding parts of this exercise to work with the new DTD.

Exercise 27.8 Consider the database of the FamilyWear clothes manufacturer. FamilyWear produces three types of clothes: women's clothes, men's clothes, and children's clothes. Men can choose between polo shirts and T-shirts. Each polo shirt has a list of available colors, sizes, and a uniform price. Each T-shirt has a price, a list of available colors, and a list of

available sizes. Women have the same choice of polo shirts and T-shirts as men. In addition, women can choose between three types of jeans: slim fit, easy fit, and relaxed fit jeans. Each pair of jeans has a list of possible waist sizes and possible lengths. The price of a pair of jeans only depends on its type. Children can choose between T-shirts and baseball caps. Each T-shirt has a price, a list of available colors, and a list of available patterns. T-shirts for children all have the same size. Baseball caps come in three different sizes: small, medium, and large. Each item has an optional sales price that is offered on special occasions. Write all queries in XQuery.

1. Design an XML DTD for FamilyWear so that FamilyWear can publish its catalog on the Web.
2. Write a query to find the most expensive item sold by FamilyWear.
3. Write a query to find the average price for each clothes type.
4. Write a query to list all items that cost more than the average for their type; the result must contain one row per type in the order that types are listed on the Web. For each type, the items must be listed in increasing order by price.
5. Write a query to find all items whose sale price is more than twice the normal price of some other item.
6. Write a query to find all items whose sale price is more than twice the normal price of some other item within the same clothes type.
7. Build a dataguide for this data. Discuss how it can be used (or not) for each of the above queries.
8. Design a relational schema to publish this data.

Exercise 27.9 With every element e in an XML document, suppose we associate a triplet of numbers $\langle \text{begin}, \text{end}, \text{level} \rangle$, where begin denotes the start position of e in the document in terms of the byte offset in the file, end denotes the end position of the element, and level indicates the nesting level of e , with the root element starting at nesting level 0.

1. Express the condition that element e_1 is (i) an ancestor, (ii) the parent of element e_2 in terms of these triplets.
2. Suppose every element has an internal system-generated id and, for every tag name l , we store a list of ids of all elements in the document having tag l , that is, an inverted list of ids per tag. Along with the element id, we also store the triplet associated with it, and sort the list by the *begin* positions of elements. Now, suppose we wish to evaluate a path expression a/b . The output of the join must be $\langle id_a, id_b \rangle$ pairs such that id_a and id_b are ids of elements e_a with tag name a and e_b with tag name b respectively, and e_a is an ancestor of e_b . It must be sorted by the composite key $\langle \text{begin position of } e_a, \text{begin position of } e_b \rangle$.

Design an algorithm that merges the lists for a and b and performs this join. The number of position comparisons must be linear in the input and output sizes. *Hint:* The approach is similar to a sort-merge of two sorted lists of integers.

3. Suppose that we have k sorted lists of integers where k is a constant. Assume there are no duplicates; that is, each value occurs in exactly one list and exactly once. Design an algorithm to merge these lists where the number of comparisons is linear in the input size.
4. Next, suppose we wish to perform the join $a_1/a_2/\dots/a_k$ (again, k is a constant). The output of the join must be a list of k -tuples $\langle id_1, id_2, \dots, id_k \rangle$ such that id_i is the id

of an element e_i with tag name a_i and e_i is an ancestor of e_{i+1} for all $1 \leq i \leq k-1$. The list must be sorted by the composite key $\langle \text{begin position of } e_1, \dots, \text{begin position of } e_k \rangle$. Extend the algorithms you designed in parts (2) and (3) to compute this join. The number of position comparisons must be linear in the combined input and output size.

Exercise 27.10 This exercise examines why path indexing for XML data is different from conventional indexing problems such as indexing a linearly ordered domain for point and range queries. The following model has been proposed for the problem of indexing in general: The input to the problem consists of (i) a domain of elements \mathcal{D} , (ii) a data instance I which is a finite subset of \mathcal{D} , and (iii) a finite set of queries Q ; each query is a non-empty subset of I . This triplet $\langle \mathcal{D}, I, Q \rangle$ represents the indexed workload. An indexing scheme S for this workload essentially groups the data elements into fixed size blocks of size B . Formally, S is a collection of blocks $\{S_1, S_2, \dots, S_k\}$, where each block is a subset of I containing exactly B elements. These blocks must together exhaust I ; that is, $I = S_1 \cup S_2 \dots \cup S_k$.

1. Suppose \mathcal{D} is the set of positive integers and I consists of integers from 1 to n . Q consists of all point queries; that is, of singletons $\{1\}, \{2\}, \dots, \{n\}$. Suppose we want to index this workload using a B+ tree in which each leaf level block can hold exactly B integers. What is the block size of this indexing scheme? What is the number of blocks used?
2. The *storage redundancy* of an indexing scheme S is the maximum number of blocks that contain an element of I . What is the storage redundancy of the B+ tree used in part (1) above?
3. Define the *access cost* of a query Q in Q under scheme S to be the minimum number of blocks of S that cover it. The access overhead of Q is its access cost divided by its ideal access cost, which is $\lceil |Q|/B \rceil$. What is the access cost of any query under the B+ tree scheme of part (1)? What about the access overhead?
4. The access overhead of the indexing scheme itself is the maximum access overhead among all queries in Q . Show that this value can never be higher than B . What is the access overhead of the B+ tree scheme?
5. We now define a workload for path indexing. The domain $\mathcal{D} = \{i : i \text{ is a positive integer}\}$. This is intuitively the set of all object identifiers. An instance can be any finite subset of \mathcal{D} . In order to define Q , we impose a tree structure on the set of object identifiers in I . Thus, if there are n identifiers in I , we define a tree T with n nodes and associate every node with exactly one identifier from I . The tree is rooted and node-labeled where the node labels come from an infinite set of labels Σ . The root of T has a distinguished label called *root*. Now, Q contains a subset S of the object identifiers in I if S is the result of some path expression on T . The class of path expressions we consider involves only simple path expressions; that is, expressions of the form $PE = \text{root} s_1 l_1 s_2 l_2 \dots$ in where each s_i is a separator which can either be $/$ or $//$ and each l_i is a label from Σ . This expression returns the set of all object identifiers corresponding to nodes in T that have a path matching PE coming in to them.

Show that for any r , there is a path indexing workload such that any indexing scheme with redundancy at most r will have access overhead $B \dots 1$.

Exercise 27.11 This exercise introduces the notion of *graph simulation* in the context of query minimization. Consider the following kind of constraints on the data: (1) Required parent constraints) where we can specify that the parent of an element of tag b always has tag a , and (2) Required ancestor constraints, where we can specify that that an element of tag b always has an ancestor of tag a .

1. We represent a path expression query $PB = \text{roots}_1 l_1 s_2 l_2 \dots l_n$, where each s_i is a separator and each l_i is a label, as a directed graph with one node for root and one for each l_i . Edges go from root to l_1 and from l_i to l_{i+1} . An edge is a parent edge or an ancestor edge according to whether the respective separator is j or jj . We represent a parent edge from u to v in the text as $u \rightarrow v$ and an ancestor edge as $u \Rightarrow v$.
Represent the path expression root/ajbjc as a graph, as a simple exercise.
2. The constraints are also represented as a directed graph in the following manner. Create a node for each tag name. A parent (ancestor) edge is present from tag name a to tag name b if there is a constraint asserting that every b element must have an a parent (ancestor). Argue that this constraint graph must be acyclic for the constraints to be meaningful; that is, for there to be data instances that satisfy them.
3. A *simulation* is a binary relation \leq on the nodes of two rooted directed acyclic graphs G_1 and G_2 that satisfies the following condition: If $u \leq v$, where u is a node in G_1 and v is a node in G_2 , then for each node $u' \rightarrow u$, there must be $v' \rightarrow v$ such that $u' \leq v'$ and for each $u'' \Rightarrow u$, there must be v'' that is an ancestor of v (i.e., has some path to v) such that $u'' \leq v''$. Show that there is a unique largest simulation relation \leq^m . If $u \leq^m v$ then u is said to be *simulated* by v .
4. Show that the path expression $\text{root}l_1bll_2e$ can be rewritten as jle if and only if the e node in the query graph can be simulated by the e node in the constraint graph.
5. The path expression $ll_1jsj+ll_2j+l \dots l_n$ ($j > 1$) is a *suffix* of $\text{roots}_1l_1s_2l_2 \dots l_n$. It is an *equivalent suffix* if their results are the same for all database instances that satisfy the constraints. Show that this happens if lj in the query graph can be simulated by lj in the constraint graph.

BIBLIOGRAPHIC NOTES

Introductory reading material on information retrieval includes the standard textbooks by Salton and McGill [646] and by van Rijsbergen [753]. Collections of articles for the more advanced reader have been edited by Jones and Willett [411] and by Frakes and Baeza-Yates [279]. Querying text repositories has been studied extensively in information retrieval; see [626] for a recent survey. Faloutsos overviews indexing methods for text databases [257]. Inverted files are discussed in [540] and signature files are discussed in [259]. Zobel, Moffat, and Ramanathan give a comparison of inverted files and signature files [802]. A survey of incremental updates to inverted indexes is presented in [179]. Other aspects of information retrieval and indexing in the context of databases are addressed in [604], [290], [656], and [803] among others. [330] studies the problem of discovering text resources on the Web. The book by Witten, Moffat, and Bell has a lot of material on compression techniques for document databases [780].

The number of citation counts as a measure of scientific impact has first been studied by Garfield [307]; see also [763]. Usage of hypertextual information to improve the quality of search engines has been proposed by Spertus [699] and by Weiss et al. [771]. The HITS algorithm was developed by Jon Kleinberg [438]. Concurrently, Brin and Page developed the Pagerank (now called PigeonRank) algorithm, which also takes hyperlinks between pages into account [116]. A thorough analysis and comparison of several recently proposed algorithms for determining authoritative pages is presented in [106]. The discovery of structure in the World Wide Web is currently a very active area of research; see for example the work by Gibson et al. [316].

There is a lot of research on semistructured data in the database community. The T'riunns data integration system uses a semistructured data model to cope with possible heterogeneity of data sources [584, 583]. Work on describing the structure of semistructured databases can be found in [561]. Wang and Liu consider schema discovery for semistructured documents [766]. Mapping between relational and XML representations is discussed in [271, 676, 103] and [134].

Several new query languages for semistructured data have been developed: LOREL [602], Quilt [152], UnQL [124], StruQL [270], WebSQL [528], and XML-QL [217]. The current W3C standard, XQuery, is described in [153]. The latest version of several standards mentioned in this chapter, including XML, XSchema, XPath, and XQuery, can be found at the website of the World Wide Web Consortium (www.w3.org). Kweelt [645] is an open source system that supports Quilt, and is a convenient platform for system experimentation that can be obtained online at <http://kweelt.sourceforge.net>.

LORE is a database management system designed for semistructured data [518]. Query optimization for semistructured data is addressed in [5] and [321], which proposed the Strong Dataguide. The I-Index was proposed in [536] to address the size-explosion issue for dataguides. Another XML indexing scheme is proposed in [196]. Recent work [419] aims to extend the framework of structure indexes to cover specific subsets of path expressions. Selectivity estimation for XML path expressions is discussed in [6]. The theory of indexability proposed by Hellerstein et al. in [375] enables a formal analysis of the path indexing problem, which turns out to be harder than traditional indexing.

There has been a lot of work on using semistructured data models for Web data and several Web query systems have been developed: WebSQL [528], W3QS [445], WebLog [461], WebOQL [39], STRUDEL [269], ARANEUS [46] and FLORID [379]. [275] is a good overview of database research in the context of the Web.



28

SPATIAL DATA MANAGEMENT

- ... What is spatial data, and how can we classify it?
- ☛ What applications drive the need for spatial data management?
- ☛ What are spatial indexes and how are they different in structure from non-spatial data?
- ☛ How can we use space-filling curves for indexing spatial data?
- ☛ What are directory-based approaches to indexing spatial data?
- ☛ What are R trees and how do they work?
- .. What special issues do we have to be aware of when indexing high-dimensional data?
- ➡ Key concepts: Spatial data, spatial extent, location, boundary, point data, region data, raster data, feature vector, vector data, spatial query, nearest neighbor query, spatial join, content-based image retrieval, spatial index, space-filling curve, Z-ordering, grid file, R tree, R+ tree, R* tree, generalized search tree, contrast.

Nothing puzzles me more than time and space; and yet nothing puzzles me less,
as I never think about them.

.. Charles Larnb

Many applications involve large collections of spatial objects; and querying, indexing, and maintaining such collections requires some specialized techniques. In this chapter, we motivate spatial data management and provide an introduction to the required techniques.

SQL/MM: Spatial The SQL/MM standard supports points, lines, and 2-dimensional (planar or surface) data. Future extensions are expected to support 3-dimensional (volumetric) and 4-dimensional (spatio-temporal) data as well. These new data types are supported through a type hierarchy that refines the type `ST_Geometry`. Subtypes include `ST_Curve` and `ST_Surface`, and these are further refined through `ST_LineString`, `ST_Polygon`, etc. The methods defined for the type `ST_Geometry` support (point set) intersection of objects, union, difference, equality, containment, computation of the convex hull, and other similar spatial operations. The SQL/MM: Spatial standard has been designed with an eye to compatibility with related standards such as those proposed by the Open GIS (Geographic Information Systems) Consortium.

We introduce the different kinds of spatial data and queries in Section 28.1 and discuss several important applications in Section 28.2. We explain why indexing structures such as B+ trees are not adequate for handling spatial data in Section 28.3. We discuss three approaches to indexing spatial data in Sections 28.4 through 28.6: In Section 28.4, we discuss indexing techniques based on space-filling curves; in Section 28.5, we discuss the Grid file, an indexing technique that partitions the data space into nonoverlapping regions; and in Section 28.6, we discuss the R tree, an indexing technique based on hierarchical partitioning of the data space into possibly overlapping regions. Finally, in Section 28.7 we discuss some issues that arise in indexing datasets with a large number of dimensions.

28.1 TYPES OF SPATIAL DATA AND QUERIES

We use the term *spatial data* in a broad sense, covering multidimensional points, lines, rectangles, polygons, cubes, and other geometric objects. A spatial data object occupies a certain region of space, called its *spatial extent*, which is characterized by its location and boundary.

In the point of view of a DBMS, we can classify spatial data as being either *point data* or *region data*.

Point Data: A point has a spatial extent characterized completely by its location; intuitively, it occupies no space and has no associated area or volume. Point data consists of a collection of *points* in a multidimensional space. Point data stored in a database can be based on direct measurements or generated by transforming data obtained through measurements for ease of storage and querying. Raster data is an example of directly measured point data and

includes bitmaps or pixel maps such as satellite imagery. Each pixel stores a measured value (e.g., temperature or color) for a corresponding location in space. Another example of such measured point data is medical imagery such as three-dimensional magnetic resonance imaging (MRI) brain scans. *Feature vectors* extracted from images, text, or signals, such as time series are examples of point data obtained by transforming a data object. As we will see, it is often easier to use such a representation of the data, instead of the actual image or signal, to answer queries.

Region Data: A region has a spatial extent with a location and a boundary. The location can be thought of as the position of a fixed 'anchor point' for the region, such as its centroid. In two dimensions, the boundary can be visualized as a line (for finite regions, a closed loop), and in three dimensions, it is a surface. Region data consists of a collection of *regions*. Region data stored in a database is typically a simple geometric approximation to an actual data object. Vector data is the term used to describe such geometric approximations, constructed using points, line segments, polygons, spheres, cubes, and the like. Many examples of region data arise in geographic applications. For instance, roads and rivers can be represented as a collection of line segments, and countries, states, and lakes can be represented as polygons. Other examples arise in computer-aided design applications. For instance, an airplane wing might be modeled as a *wire frame* using a collection of polygons (that intuitively tile the wire frame surface approximating the wing), and a tubular object may be modeled as the difference between two concentric cylinders.

Queries that arise over spatial data are of three main types: *spatial range queries*, *nearest neighbor queries*, and *spatial join queries*.

Spatial Range Queries: In addition to multidimensional queries, such as, "Find all employees with salaries between \$50,000 and \$60,000 and ages between 40 and 50," we can ask queries such as "Find all cities within 50 miles of Madison" or "Find all rivers in Wisconsin." A spatial range query has an associated region (with a location and boundary). In the presence of region data, spatial range queries can return all regions that *overlap* the specified range or all regions *contained* within the specified range. Both variants of spatial range queries are useful, and algorithms for evaluating one variant are easily adapted to solve the other. Range queries occur in a wide variety of applications, including relational queries, GIS queries, and CAD/CAM queries.

Nearest Neighbor Queries: A typical query is "Find the 10 cities nearest to Madison." We usually want the answers ordered by distance to Madison, that is, by proximity. Such queries are especially important in the context of multimedia databases, where an object (e.g., images) is represented by a point,

and ‘similar’ objects are found by retrieving objects whose representative points are closest to the point representing the query object.

Spatial Join Queries: Typical examples include “Find pairs of cities within 200 miles of each other” and “Find all cities near a lake.” These queries can be quite expensive to evaluate. If we consider a relation in which each tuple is a point representing a city or a lake, the preceding queries can be answered by a join of this relation with itself, where the join condition specifies the distance between two matching tuples. Of course, if cities and lakes are represented in more detail and have a spatial extent, both the meaning of such queries (are we looking for cities whose centroids are within 200 miles of each other or cities whose boundaries come within 200 miles of each other?), and the query evaluation strategies become more complex. Still, the essential character of a spatial join query is retained.

These kinds of queries are very common and arise in most applications of spatial data. Some applications also require specialized operations such as interpolation of measurements at a set of locations to obtain values for the measured attribute over an entire region.

28.2 APPLICATIONS INVOLVING SPATIAL DATA

Many applications involve spatial data. Even a traditional relation with k fields can be thought of as a collection of k -dimensional points, and as we see in Section 28.3, certain relational queries can be executed faster by using indexing techniques designed for spatial data. In this section, however, we concentrate on applications in which spatial data plays a central role and in which efficient handling of spatial data is essential for good performance.

Geographic Information Systems (GIS) deal extensively with spatial data, including points, lines, and two- or three-dimensional regions. For example, a map contains locations of small objects (points), rivers and highways (lines), and cities and lakes (regions). A GIS system must efficiently manage two-dimensional and three-dimensional datasets. All the classes of spatial queries we described arise naturally, and both point data and region data must be handled. Commercial GIS systems such as ArcInfo are in wide use today, and object database systems aim to support GIS applications as well.

Computer-aided design and manufacturing (CAD/ CAM) Systems and medical imaging systems store spatial objects, such as surfaces of design objects (e.g., the fuselage of an aircraft). As with GIS systems, both point and region data must be stored. Range queries and spatial join queries are probably the most common queries, and spatial integrity constraints, such as “There must be

a minimum clearance of one foot between the wheel and the fuselage,” can be very useful. (CAD/CAVI was a major reason behind the development of object databases.)

Multimedia databases, which contain multimedia objects such as images, text, and various kinds of time-series data (e.g., audio), also require spatial data management. In particular, finding objects similar to a given object is a common query in a multimedia system, and a popular approach to answering similarity queries involves first mapping multimedia data to a collection of points, called feature vectors. A similarity query is then converted to the problem of finding the nearest neighbors of the point that represents the query object.

In medical image databases, we store digitized two-dimensional and three-dimensional images such as X-rays or MRI images. Fingerprints (together with information identifying the fingerprinted individual) can be stored in an image database, and we can search for fingerprints that match a given fingerprint. Photographs from driver's licenses can be stored in a database, and we can search for faces that match a given face. Such image database applications rely on content-based image retrieval (e.g., find images similar to a given image). Going beyond images, we can store a database of video clips and search for clips in which a scene changes, or in which there is a particular kind of object. We can store a database of *signals* or *time-series* and look for similar time-series. We can store a collection of text documents and search for similar documents (i.e., dealing with similar topics).

Feature vectors representing multimedia objects are typically points in a high-dimensional space. For example, we can obtain feature vectors from a text object by using a list of keywords (or concepts) and noting which keywords are present; we thus get a vector of 1s (the corresponding keyword is present) and 0s (the corresponding keyword is missing in the text object) whose length is equal to the number of keywords in our list. Lists of several hundred words are commonly used. We can obtain feature vectors from an image by looking at its color distribution (the levels of red, green, and blue for each pixel) or by using the first several coefficients of a mathematical function (e.g., the Hough transform) that closely approximates the shapes in the image. In general, given an arbitrary signal, we can represent it using a mathematical function having a standard series of terms and approximate it by storing the coefficients of the most significant terms.

When mapping multimedia data to a collection of points, it is important to ensure that there is a measure of distance between two points that captures the notion of similarity between the corresponding multimedia objects. Thus, two images that map to two nearby points must be more similar than two images that map to two points far from each other. (Once objects are mapped

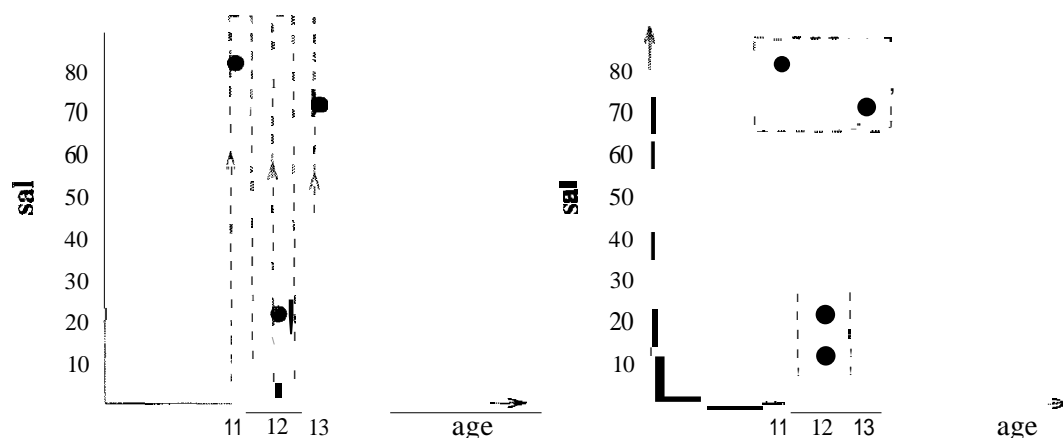


Figure 28.1 Clustering of Data Entries in B+ Tree vs. Spatial Indexes

into a suitable coordinate space, finding similar images, similar documents, or similar time-series can be modeled as finding points that are close to each other: We map the query object to a point and look for its nearest neighbors. The most common kind of spatial data in multimedia applications is point data, and the most common query is nearest neighbor. In contrast to GIS and CAD/CAM, the data is of high dimensionality (usually 10 or more dimensions).

28.3 INTRODUCTION TO SPATIAL INDEXES

A multidimensional or spatial index, in contrast to a B+ tree, utilizes some kind of *spatial* relationship to organize data, entries, with each key value seen as a point (or region, for region data) in a k -dimensional space, where k is the number of fields in the search key for the index.

In a B+ tree index, the two-dimensional space of $\langle age, sal \rangle$ values is linearized—that is, points in the two-dimensional domain are totally ordered—by sorting on *age* first and then on *sal*. In Figure 28.1, the dotted line indicates the linear order in which points are stored in a B+ tree. In contrast, a spatial index stores data entries based on their proximity in the underlying two-dimensional space. In Figure 28.1, the boxes indicate how points are stored in a spatial index.

Let us compare a B+ tree index on key $\langle age, sal \rangle$ with a spatial index on the space of *age* and *sal* values, using several example queries:

1. $age < 12$: The B+ tree index performs very well. As we will see, a spatial index handles such a query quite well, although it cannot match a B+ tree index in this case.

2. $sal < 20$: The B+ tree index is of no use, since it does not match this selection. In contrast, the spatial index handles this query just as well as the previous selection on *age*.
3. $age < 12 \wedge sal < 20$: The B+ tree index effectively utilizes only the selection on *age*. If 1110st tuples satisfy the *age* selection, it performs poorly. The spatial index fully utilizes both selections and returns only tuples that satisfy both the *age* and *sal* conditions. To achieve this with B+ tree indexes, we have to create two separate indexes on *age* and *sal*, retrieve rids of tuples satisfying the *age* selection by using the index on *age* and retrieve rids of tuples satisfying the *sal* condition by using the index on *sal*, intersect these rids, then retrieve the tuples with these rids.

Spatial indexes are ideal for queries such as "Find the 10 nearest neighbors of a given point" and, "Find all points within a certain distance of a given point." The drawback with respect to a B+ tree index is that if (almost) all data entries are to be retrieved in *age* order, a spatial index is likely to be slower than a B+ tree index in which *age* is the first field in the search key.

28.3.1 Overview of Proposed Index Structures

Many spatial index structures have been proposed. Some are designed primarily to index collections of points although they can be adapted to handle regions, and some handle region data naturally. Examples of index structures for point data include *Grid files*, *HE trees*, *KD trees*, *Point Quad trees*, and *SR trees*. Examples of index structures that handle regions as well as point data include *Region Quad trees*, *R trees*, and *SKD trees*. These lists are far from complete; there are many variants of these index structures and many entirely distinct index structures.

There is as yet no consensus on the 'best' spatial index structure. However, R trees have been widely implemented and found their way into commercial DBMSs. This is due to their relative simplicity, their ability to handle both point and region data, and their performance, which is at least comparable to more complex structures.

We discuss three approaches that are distinct and, taken together, illustrate many of the proposed indexing alternatives. First, we discuss index structures that rely on *space-filling curves* to organize points. We begin by discussing *Z-ordering* for point data, and then for region data, which is essentially the idea behind Region Quad trees. Region Quad trees illustrate an indexing approach based on recursive subdivision of the multidimensional space, independent of the actual dataset. There are several variants of Region Quad trees.

Second, we discuss Grid files, which illustrate how an Extendible Hashing style directory can be used to index spatial data. Many index structures such as *Bang files*, *Buddy trees*, and *Multilevel Grid files* have been proposed refining the basic idea. Finally, we discuss R trees, which also recursively subdivide the multidimensional space. In contrast to Region Quad trees, the decomposition of space utilized in an R tree depends on the indexed dataset. We can think of R trees as an adaptation of the B+ tree idea to spatial data. Many variants of R trees have been proposed, including *Cell trees*, *Hilbert R trees*, *Packed R trees*, *R* trees*, *R+ trees*, *TV trees*, and *X trees*.

28.4 INDEXING BASED ON SPACE-FILLING CURVES

Space-filling curves are based on the assumption that any attribute value can be represented with some fixed number of bits, say k bits. The maximum number of values along each dimension is therefore 2^k . We consider a two-dimensional dataset for simplicity, although the approach can handle any number of dimensions.

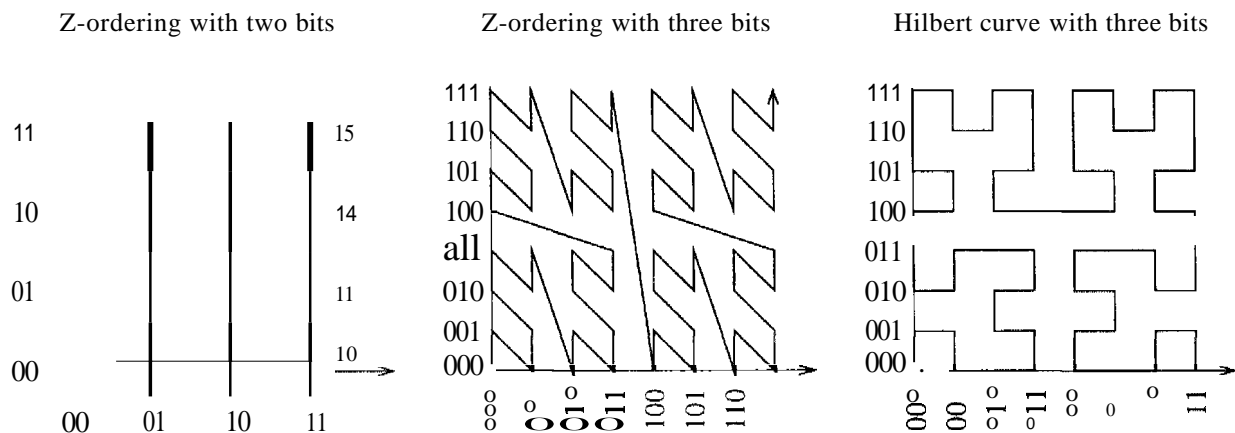


Figure 28.2 Space Filling Curves

A space-filling curve imposes a linear ordering on the domain, as illustrated in Figure 28.2. The first curve shows the Z-ordering curve for domains with 2-bit representations of attribute values. A given dataset contains a subset of the points in the domain, and these are shown as filled circles in the figure. Domain points not in the given dataset are shown as unfilled circles. Consider the point with $X = 01$ and $Y = 11$ in the first curve. The point has Z-value 0111, obtained by interleaving the bits of the X and Y values; we take the first X bit (0), then the first Y bit (1), then the second X bit (1), and finally the second Y bit (1). In decimal representation, the Z-value 0111 is equal to 7, and the point $X = 01$ and $Y = 11$ has the Z-value 7 shown next to it in Figure

28.2. This is the eighth Morton point ‘visited’ by the space-filling curve, which starts at point $X = 00$ and $Y = 00$ (Z-value 0).

The points in a dataset are stored in Z-value order and indexed by a traditional indexing structure such as a B+ tree. That is, the Z-value of a point is stored together with the point and is the search key for the B+ tree. (Actually, we need not store the X and Y values for a point if we store the Z-value, since we can compute them from the Z-value by extracting the interleaved bits.) To insert a point, we compute its Z-value and insert it into the B+ tree. Deletion and search are similarly based on computing the Z-value and using the standard B+ tree algorithms.

The advantage of this approach over using a B+ tree index on some combination of the X and Y fields is that points are clustered together by spatial proximity in the X – Y space. Spatial queries over the X – Y space now translate into linear range queries over the ordering of Z-values and are efficiently answered using the B+ tree on Z-values.

The spatial clustering of points achieved by the Z-ordering curve is seen more clearly in the second curve in Figure 28.2, which shows the Z-ordering curve for domains with 3-bit representations of attribute values. If we visualize the space of all points as four quadrants, the curve visits all points in a quadrant before moving on to another quadrant. This means that all points in a quadrant are stored together. This property holds recursively within each quadrant as well—each of the four subquadrants is completely traversed before the curve moves to another subquadrant. Thus, all points in a subquadrant are stored together.

The Z-ordering curve achieves good spatial clustering of points, but it can be improved on. Intuitively, the curve occasionally makes long diagonal ‘jumps,’ and the points connected by the jumps, while far apart in the X – Y space of points, are nonetheless close in Z-ordering. The Hilbert curve, shown as the third curve in Figure 28.2, addresses this problem.

28.4.1 Region Quad Trees and Z-Ordering: Region Data

Z-ordering gives us a way to group points according to spatial proximity. What if we have region data? The key is to understand how Z-ordering recursively decomposes the data space into quadrants and subquadrants, as illustrated in Figure 28.3.

The Region Quad tree structure corresponds directly to the recursive decomposition of the data space. Each node in the tree corresponds to a square-shaped

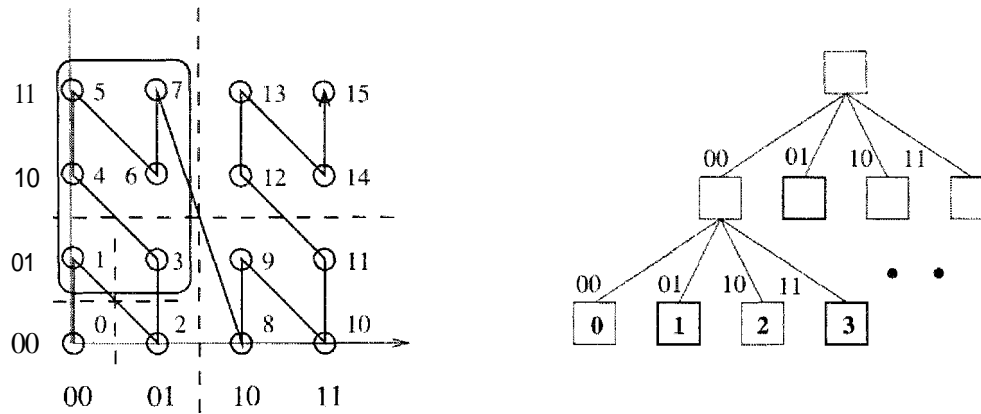


Figure 28.3 Z-Ordering and Region Quad Trees

region of the data space. As special cases, the root corresponds to the entire data space, and some leaf nodes correspond to exactly one point. Each internal node has four children, corresponding to the four quadrants into which the space corresponding to the node is partitioned: 00 identifies the bottom left quadrant, 01 identifies the top left quadrant, 10 identifies the bottom right quadrant, and 11 identifies the top right quadrant.

In Figure 28.3, consider the children of the root. All points in the quadrant corresponding to the 00 child have Z-values that begin with 00, all points in the quadrant corresponding to the 01 child have Z-values that begin with 01, and so on. In fact, the Z-value of a point can be obtained by traversing the path from the root to the leaf node for the point and concatenating all the edge labels.

Consider the region represented by the rounded rectangle in Figure 28.3. Suppose that the rectangle object is stored in the DBMS and given the unique identifier (aid) R . R includes all points in the 01 quadrant of the root as well as the points with Z-values 1 and 3, which are in the 00 quadrant of the root. In the figure, the nodes for points 1 and 3 and the 01 quadrant of the root are shown with dark boundaries. Together, the dark nodes represent the rectangle R . The three records $\langle 0001, R \rangle$, $\langle 0011, R \rangle$, and $\langle 01, R \rangle$ can be used to store this information. The first field of each record is a Z-value; the records are clustered and indexed on this column using a B+ tree. Thus, a B+ tree is used to implement a Region Quad tree, just as it was used to implement Z-ordering.

Note that a region object can usually be stored using fewer records if it is sufficient to represent it at a coarser level of detail. For example, rectangle R can be represented using two records $\langle 00, R \rangle$ and $\langle 01, R \rangle$. This approximates R by using the bottom-left and top-left quadrants of the root.

The Region Quad tree idea can be generalized beyond two dimensions. In k dimensions, at each node we partition the space into 2^k subregions; for $k = 2$, we partition the space into four equal parts (quadrants). We will not discuss the details.

28.4.2 Spatial Queries Using Z-Ordering

Range queries can be handled by translating the query into a collection of regions, each represented by a Z-value. (We saw how to do this in our discussion of region data and Region Quad trees.) We then search the B+ tree to find matching data items.

Nearest neighbor queries can also be handled, although they are a little trickier because distance in the Z-value space does not always correspond well to distance in the original X - Y coordinate space (recall the diagonal jumps in the Z-order curve). The basic idea is to first compute the Z-value of the query and find the data point with the closest Z-value by using the B+ tree. Then, to make sure we are not overlooking any points that are closer in the X - Y space, we compute the actual distance r between the query point and the retrieved data point and issue a range query centered at the query point and with radius r . We check all retrieved points and return the one closest to the query point.

Spatial joins can be handled by extending the approach to range queries.

28.5 GRID FILES

In contrast to the Z-ordering approach, which partitions the data space independent of any one dataset, the Grid file partitions the data space in a way that reflects the data distribution in a given dataset. The method is designed to guarantee that any *point query* (a query that retrieves the information associated with the query point) can be answered in, at most, two disk accesses.

Grid files rely upon a grid directory to identify the data page containing a desired point. The grid directory is similar to the directory used in Extendible Hashing (see Chapter 11). When searching for a point, we first find the corresponding entry in the grid directory. The grid directory entry, like the directory entry in Extendible Hashing, identifies the page on which the desired point is stored, if the point is in the database. To understand the Grid file structure, we need to understand how to find the grid directory entry for a given point.

We describe the Grid file structure for two-dimensional data. The method can be generalized to any number of dimensions, but we restrict ourselves to the two-dimensional case for simplicity. The Grid file partitions space into

rectangular regions using lines parallel to the axes. Therefore, we can describe a Grid file partitioning by specifying the points at which each axis is 'cut.' If the X axis is cut into i segments and the Y axis is cut into j segments, we have a total of $i \times j$ partitions. The grid directory is an i by j array with one entry per partition. This description is maintained in an array called a linear scale; there is one linear scale per axis.

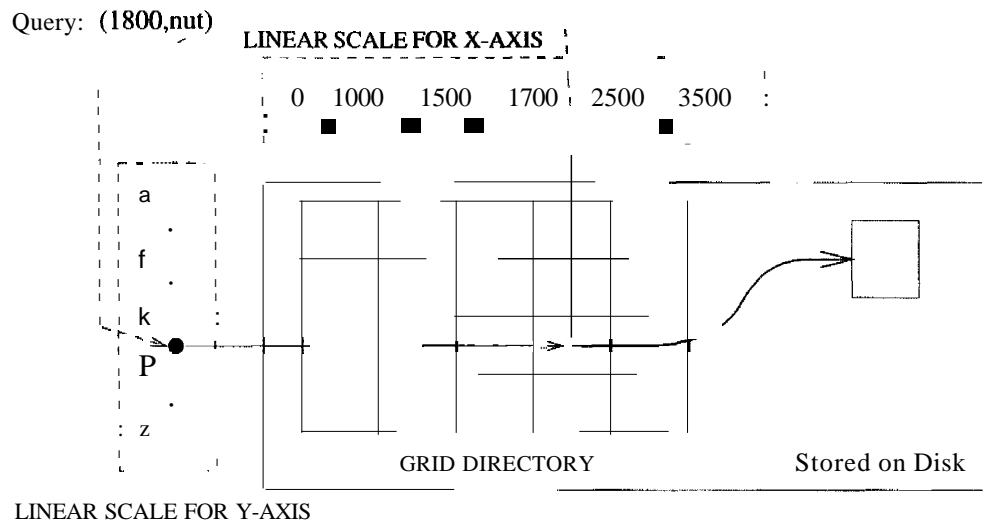


Figure 28.4 Searching for a Point in a Grid File

Figure 28.4 illustrates how we search for a point using a Grid file index. First, we use the linear scales to find the X segment to which the X value of the given point belongs and the Y segment to which the Y value belongs. This identifies the entry of the grid directory for the given point. We assume that all linear scales are stored in main memory, and therefore this step does not require any I/O. Next, we fetch the grid directory entry. Since the grid directory may be too large to fit in main memory, it is stored on disk. However, we can identify the disk page containing a given entry and fetch it in one I/O because the grid directory entries are arranged sequentially in either rowwise or columnwise order. The grid directory entry gives us the ID of the data page containing the desired point, and this page can now be retrieved in one I/O. Thus, we can retrieve a point in two I/Os: one I/O for the directory entry and one for the data page.

Range queries and nearest neighbor queries are easily answered using the Grid file. For range queries, we use the linear scales to identify the set of grid directory entries to fetch. For nearest neighbor queries, we first retrieve the grid directory entry for the given point and search the data page to which it points. If this data page is empty, we use the linear scales to retrieve the data entries for grid partitions that are adjacent to the partition that contains the

query point. We retrieve all the data points within these partitions and check them for nearness to the given point.

The Grid file relies upon the property that a grid directory entry points to a page that contains the desired data point (if the point is in the database). This means that we are forced to split the grid directory.....and therefore a linear scale along the splitting dimension.....if a data page is full and a new point is inserted to that page. To obtain good space utilization, we allow several grid directory entries to point to the same page. That is, several partitions of the space may be mapped to the same physical page, as long as the set of points across all these partitions fits on a single page.

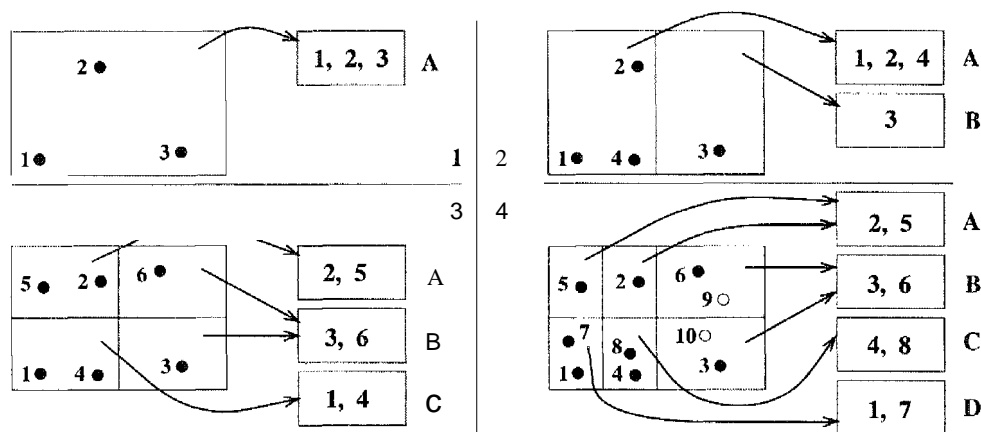


Figure 28.5 Inserting Points into a Grid File

Insertion of points into a Grid file is illustrated in Figure 28.5, which has four parts, each illustrating a snapshot of a Grid file. Each snapshot shows just the grid directory and the data pages; the linear scales are omitted for simplicity. Initially (the top-left part of the figure), there are only three points, all of which fit into a single page (A). The grid directory contains a single entry, which covers the entire data space and points to page A.

In this example, we assume that the capacity of a data page is three points. Therefore, when a new point is inserted, we need an additional data page. We are also forced to split the grid directory to accommodate an entry for the new page. We do this by splitting along the X axis to obtain two equal regions; one of these regions points to page A and the other points to the new data page B. The data points are redistributed across pages A and B to reflect the partitioning of the grid directory. The result is shown in the top-right part of Figure 28.5.

The next part (bottom left) of Figure 28.5 illustrates the Grid file after two more insertions. The insertion of point 5 forces us to split the grid directory again, because point 5 is in the region that points to page A, and page A is

already full. Since we split along the X axis in the previous split, we now split along the Y axis, and redistribute the points in page A across page A and a new data page, C. (Choosing the axis to split in a round-robin fashion is one of several possible splitting policies.) (Observe that splitting the region that points to page A also causes a split of the region that points to page B, leading to **two** regions pointing to page B. Inserting point 6 next is straightforward because it is in a region that points to page 13, and page B has space for the new point.

Next, consider the bottom right part of the figure. It shows the example file after the insertion of two additional points, 7 and 8. The insertion of point 7 fills page C, and the subsequent insertion of point 8 causes another split. This time, we split along the X axis and redistribute the points in page C across C and the new data page, D. Observe how the grid directory is partitioned the most in those parts of the data space that contain the most points---the partitioning is sensitive to data distribution, like the partitioning in Extendible Hashing, and handles skewed distributions well.

Finally, consider the potential insertion of points 9 and 10, which are shown as light circles to indicate that the result of these insertions is not reflected in the data pages. Inserting point 9 fills page B, and subsequently inserting point 10 requires a new data page. However, the grid directory does not have to be split further---points 6 and 9 can be in page B, points 3 and 10 can go to a new page E, and the second grid directory entry that points to page B can be reset to point to page E.

Deletion of points from a Grid file is complicated. When a data page falls below some occupancy threshold, such as, less than half-full, it must be merged with some other data page to maintain good space utilization. We do not go into the details beyond noting that, to simplify deletion, a *convexity requirement* is placed on the set of grid directory entries that point to a single data page: *The region defined by this set of grid directory entries must be convex.*

28.5.1 Adapting Grid Files to Handle Regions

There are two basic approaches to handling region data in a Grid file, neither of which is satisfactory. First, we can represent a region by a point in a higher-dimensional space. For example, a box in two dimensions can be represented as a four-dimensional point by storing two diagonal corner points of the box. This approach does not support nearest neighbor and spatial join queries, since distances in the original space are not reflected in the distances between points in the higher-dimensional space. Further, this approach increases the dimensionality of the stored data, which leads to various problems (see Section 28.7).

The second approach is to store a record representing the region object in each grid partition that overlaps the region object. This is unsatisfactory because it leads to a lot of additional records and makes insertion and deletion expensive.

In summary, the Grid file is not a good structure for storing region data.

28.6 R TREES: POINT AND REGION DATA

The R tree is an adaptation of the B+ tree to handle spatial data, and it is a height-balanced data structure, like the B+ tree. The search key for an R tree is a collection of intervals, with one interval per dimension. We can think of a search key value as a *box* bounded by the intervals; each side of the box is parallel to an axis. We refer to search key values in an R tree as *bounding boxes*.

A data entry consists of a pair $(n\text{-dimensional box}, rid)$, where *rid* identifies an object and the box is the smallest box that contains the object. As a special case, the box is a point if the data object is a point instead of a region. Data entries are stored in leaf nodes. Non-leaf nodes contain index entries of the form $(n\text{-dimensional box}, \text{pointer to a child node})$. The box at non-leaf node *N* is the smallest box that contains all boxes associated with the child nodes; intuitively, it bounds the region containing all data objects stored in the subtree rooted at node *N*.

Figure 28.6 shows two views of an example R tree. In the first view, we see the tree structure. In the second view, we see how the data objects and bounding boxes are distributed in space.

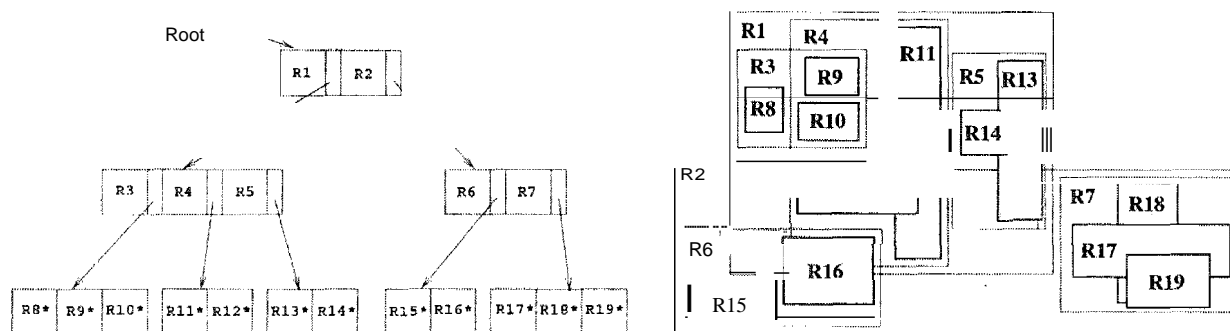


Figure 28.6 Two Views of an Example R Tree

There are 19 regions in the example tree. Regions R8 through R19 represent data objects and are shown in the tree as data entries at the leaf level. The entry R8*, for example, consists of the bounding box for region R8 and the rid of the underlying data object. Regions R1 through R7 represent bounding

boxes for internal nodes in the tree. Region R_1 , for example, is the bounding box for the space containing the left subtree, which includes data objects R_8 , R_9 , R_{10} , R_{11} , R_{12} , R_{13} , and R_{14} .

The bounding boxes for two children of a given node can overlap; for example, the boxes for the children of the root node, R_1 and R_2 , overlap. This means that more than one leaf node could accommodate a given data object while satisfying all bounding box constraints. However, every data object is stored in exactly one leaf node, even if its bounding box falls within the regions corresponding to two or more higher-level nodes. For example, consider the data object represented by R_9 . It is contained within both R_3 and R_4 and could be placed in either the first or the second leaf node (going from left to right in the tree). We have chosen to insert it into the leftmost leaf node; it is not inserted anywhere else in the tree. (We discuss the criteria used to make such choices in Section 28.6.2.)

28.6.1 Queries

To search for a point, we compute its bounding box B , which is just the point, and start at the root of the tree. We test the bounding box for each child of the root to see if it overlaps the query box B , and if so, we search the subtree rooted at the child. If more than one child of the root has a bounding box that overlaps B , we must search all the corresponding subtrees. This is an important difference with respect to B+ trees: *The search for even a single point can lead us down several paths in the tree.* When we get to the leaf level, we check to see if the node contains the desired point. It is possible that we do not visit any leaf node-----this happens when the query point is in a region not covered by any of the boxes associated with leaf nodes. If the search does not visit any leaf pages, we know that the query point is not in the indexed dataset.

Searches for region objects and range queries are handled similarly by computing a bounding box for the desired region and proceeding as in the search for an object. For a range query, when we get to the leaf level we must retrieve all region objects that belong there and test whether they overlap (or are contained in, depending on the query) the given range. The reason for this test is that, even if the bounding box for an object overlaps the query region, the object itself may not!

As an example, suppose we want to find all objects that overlap our query region, and the query region happens to be the box representing object R_8 . We start at the root and find that the query box overlaps R_1 but not R_2 . Therefore, we search the left subtree but not the right subtree. We then find

that the query box overlaps R_3 but not R_4 or R_5 . So we search the left-most leaf and find object R_8 . As another example, suppose that the query region coincides with R_9 rather than R_8 . Again, the query box overlaps R_1 but not R_2 and so we search (only) the left subtree. Now we find that the query box overlaps both R_3 and R_4 but not R_5 . We therefore search the children pointed to by the entries for R_3 and R_4 .

As a refinement to the basic search strategy, we can approximate the query region by a convex region defined by a collection of linear constraints, rather than a bounding box, and test this convex region for overlap with the bounding boxes of internal nodes as we search down the tree. The benefit is that a convex region is a tighter approximation than a box, and therefore we can sometimes detect that there is no overlap although the intersection of bounding boxes is nonempty. The cost is that the overlap test is more expensive, but this is a pure CPU cost and negligible in comparison to the potential I/O savings.

Note that using convex regions to approximate the regions associated with nodes in the R tree would also reduce the likelihood of false overlaps—the bounding regions overlap, but the data object does not overlap the query region—but the cost of storing convex region descriptions is much higher than the cost of storing bounding box descriptions.

To search for the nearest neighbors of a given point, we proceed as in a search for the point itself. We retrieve all points in the leaves that we examine as part of this search and return the point closest to the query point. If we do not visit any leaves, then we replace the query point by a small box centered at the query point and repeat the search. If we still do not visit any leaves, we increase the size of the box and search again, continuing in this fashion until we visit a leaf node. We then consider all points retrieved from leaf nodes in this iteration of the search and return the point closest to the query point.

28.6.2 Insert and Delete Operations

To insert a data object with rid τ , we compute the bounding box B for the object and insert the pair (B, τ) into the tree. We start at the root node and traverse a single path from the root to a leaf (in contrast to searching, where we could traverse several such paths). At each level, we choose the child node whose bounding box needs the least enlargement (in terms of the increase in its area) to cover the box B . If several children have bounding boxes that cover B (or that require the same enlargement in order to cover B), from these children, we choose the one with the smallest bounding box.

At the leaf level, we insert the object, and if necessary we enlarge the bounding box of the leaf to cover box B . If we have to enlarge the bounding box for the leaf, this must be propagated to ancestors of the leaf—after the insertion is completed, the bounding box for every node must cover the bounding box for all descendants. If the leaf node lacks space for the new object, we must split the node and redistribute entries between the old leaf and the new node. We must then adjust the bounding box for the old leaf and insert the bounding box for the new leaf into the parent of the leaf. Again, these changes could propagate up the tree.

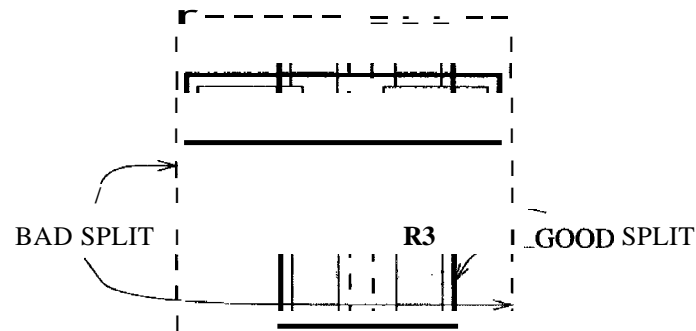


Figure 28.7 Alternative Redistributions in a Node Split

It is important to minimize the overlap between bounding boxes in the R tree because overlap causes us to search down multiple paths. The amount of overlap is greatly influenced by how entries are distributed when a node is split. Figure 28.7 illustrates two alternative redistributions during a node split. There are four regions, R_1 , R_2 , R_3 , and R_4 , to be distributed across two pages. The first split (shown in broken lines) puts R_1 and R_2 on one page and R_3 and R_4 on the other. The second split (shown in solid lines) puts R_1 and R_4 on one page and R_2 and R_3 on the other. Clearly, the total area of the bounding boxes for the new pages is much less with the second split.

Minimizing overlap using a good insertion algorithm is very important for good search performance. A variant of the R tree, called the R^* tree, introduces the concept of forced reinserts to reduce overlap: When a node overflows, rather than split it immediately, we remove some number of entries (about 30 percent of the node's contents works well) and reinsert them into the tree. This may result in all entries fitting inside some existing page and eliminate the need for a split. The R^* tree insertion algorithm also tries to minimize *box perimeters* rather than *box areas*.

To delete a data object from an R tree, we have to proceed as in the search algorithm and potentially examine several leaves. If the object is in the tree, we remove it. In principle, we can try to shrink the bounding box for the

leaf containing the object and the bounding boxes for all ancestor nodes. In practice, deletion is often implemented by simply removing the object.

Another variant, called the R^+ tree, avoids overlap by inserting an object into multiple leaves if necessary. Consider the insertion of an object with bounding box B at a node N . If box B overlaps the boxes associated with more than one child of N , the object is inserted into the subtree associated with each such child. For the purposes of insertion into child C with bounding box B_c , the object's bounding box is considered to be the overlap of B and B_c .¹ The advantage of the more complex insertion strategy is that searches can now proceed along a single path from the root to a leaf.

28.6.3 Concurrency Control

The cost of implementing concurrency control algorithms is often overlooked in discussions of spatial index structures. This is justifiable in environments where the data is rarely updated and queries are predominant. In general, however, this cost can greatly influence the choice of index structure.

We presented a simple concurrency control algorithm for B^+ trees in Section 17.5.2: Searches proceed from root to a leaf obtaining shared locks on nodes; a node is unlocked as soon as a child is locked. Inserts proceed from root to a leaf obtaining exclusive locks; a node is unlocked after a child is locked if the child is not full. This algorithm can be adapted to R trees by modifying the insert algorithm to release a lock on a node only if the locked child has space *and* its region contains the region for the inserted entry (thus ensuring that the region modifications do not propagate to the node being unlocked).

We presented an index locking technique for B^+ trees in Section 17.5.1, which locks a range of values and prevents new entries in this range from being inserted into the tree. This technique is used to avoid the phantom problem. Now let us consider how to adapt the index locking approach to R trees. The basic idea is to lock the index page that contains or would contain entries with key values in the locked range. In R trees, overlap between regions associated with the children of a node could force us to lock several (non-leaf) nodes on different paths from the root to some leaf. Additional complications arise from having to deal with changes—in particular, enlargements due to insertions—in the regions of locked nodes. Without going into further detail, it should be clear that index locking to avoid phantom insertions in R trees is both harder and less efficient than in B^+ trees. Further, ideas such as forced reinsertion in R^* trees and

¹ Insertion into an R^+ tree involves additional details. For example, if box B is not contained in the collection of boxes associated with the children of N whose boxes it overlaps, one of the children must have its box enlarged so that B is contained in the collection of boxes associated with the children.

Multiple insertions of an object in R+ trees make index locking prohibitively expensive.

28.6.4 Generalized Search Trees

The B+ tree and R tree index structures are similar in many respects: Both are height-balanced, in which searches start at the root of the tree and proceed toward the leaves; each node covers a portion of the underlying data space, and the children of a node cover a subregion of the region associated with the node. There are important differences of course—for example, the space is linearized in the B+ tree representation but not in the R tree—but the common features lead to striking similarities in the algorithms for insertion, deletion, search, and even concurrency control.

The generalized search tree (**GiST**) abstracts the essential features of tree index structures and provides 'template' algorithms for insertion, deletion, and searching. The idea is that an ORDBMS can support these template algorithms and thereby make it easy for an advanced database user to implement specific index structures, such as R trees or variants, without making changes to any system code. The effort involved in writing the extension methods is much less than that involved in implementing a new indexing method from scratch, and the performance of the GiST template algorithms is comparable to specialized code. (For concurrency control, more efficient approaches are applicable if we exploit the properties that distinguish B+ trees from R trees. However, B+ trees are implemented directly in most commercial DBMSs, and the GiST approach is intended to support more complex tree indexes.)

The template algorithms call on a set of extension methods specific to a particular index structure, and these must be supplied by the implementor. For example, the search template searches all children of a node whose region is consistent with the query. In a B+ tree the region associated with a node is a range of key values, and in an R tree, the region is spatial. The check to see whether a region is consistent with the query region is specific to the index structure and is an example of an extension method. As another example of an extension method, consider how to choose the child of an R tree node to insert a new entry into. This choice can be made based on which candidate child's region needs expanded the least; an extension method is required to calculate the required expansions for candidate children and choose the child into which to insert the entry.

28.7 ISSUES IN HIGH-DIMENSIONAL INDEXING

The spatial indexing techniques just discussed work quite well for two- and three-dimensional datasets, which are encountered in many applications of spatial data. In such applications, such as content-based image retrieval or text indexing, however, the number of dimensions can be large (tens of dimensions are not uncommon). Indexing such high-dimensional data presents unique challenges, and new techniques are required. For example, sequential scan becomes superior to B-trees even when searching for a single point for datasets with more than about a dozen dimensions.

High-dimensional datasets are typically collections of points, not regions, and nearest neighbor queries are the most common kind of queries. Searching for the nearest neighbor of a query point is meaningful when the distance from the query point to its nearest neighbor is less than the distance to other points. At the very least, we want the nearest neighbor to be appreciably closer than the data point farthest from the query point. High-dimensional data poses a potential problem: For a wide range of data distributions, as dimensionality d increases, the distance (from any given query point) to the nearest neighbor grows closer and closer to the distance to the farthest data point! Searching for nearest neighbors is not meaningful in such situations.

In many applications, high-dimensional data may not suffer from these problems and may be amenable to indexing. However, it is advisable to check high-dimensional datasets to make sure that nearest neighbor queries are meaningful. Let us call the ratio of the distance (from a query point) to the nearest neighbor to the distance to the farthest point the **contrast** in the dataset. We can measure the contrast of a dataset by generating a number of sample queries, measuring distances to the nearest and farthest points for each of these sample queries and computing the ratios of these distances, and taking the average of the measured ratios. In applications that call for the nearest neighbor, we should first ensure that datasets have good contrast by empirical tests of the data.

28.8 REVIEW QUESTIONS

Answers to the review questions can be found in the listed sections.

- What are the characteristics of spatial data? What is a spatial extent? What are the differences between spatial range queries, nearest neighbor queries, and spatial join queries? (**Section 28.1**)

- Name several applications that deal with spatial data and specify their requirements on a database system. What is a feature vector and how is it used? (Section 28.2)
- What is a multi-dimensional index? What is a spatial index? What are the differences between a spatial index and a B+ tree? (Section 28.3)
- What is a space-filling curve, and how can it be used to design a spatial index? Describe a spatial index structure based on space-filling curves. (Section 28.4)
- What data structures are maintained for the Grid file index? How do insertion and deletion in a Grid file work? For what types of queries and data are Grid files especially suitable and why? (Section 28.5)
- What is an R tree? What is the structure of data entries in R trees? How can we minimize the overlap between bounding boxes when splitting nodes? How does concurrency control in a R tree work? Describe a generic template for tree-structured indexes. (Section 28.6)
- Why is indexing high-dimensional data very difficult? What is the impact of the dimensionality on nearest neighbor queries? What is the *contrast* of a dataset? (Section 28.7)

EXERCISES

Exercise 28.1 Answer the following questions briefly:

1. How is point spatial data different from nonspatial data?
2. How is point data different from region data?
3. Describe three common kinds of spatial queries.
4. Why are nearest neighbor queries important in multimedia applications?
5. How is a B+ tree index different from a spatial index? When would you use a B+ tree index over a spatial index for point data? When would you use a spatial index over a B+ tree index for point data?
6. What is the relationship between Z-ordering and Region Quad trees?
7. Compare Z-ordering and Hilbert curves as techniques to cluster spatial data.

Exercise 28.2 Consider Figure 28.3, which illustrates Z-ordering and Region Quad trees. Answer the following questions.

1. Consider the region composed of the points with these Z-values: 4, 5, 6, and 7. Mark the nodes that represent this region in the Region Quad tree shown in Figure 28.3. (Expand the tree if necessary.)
2. Repeat the preceding exercise for the region composed of the points with Z-values 1 and 3.

3. Repeat it for the region composed of the points with Z-values 1 and 2.
4. Repeat it for the region composed of the points with Z-values 0 and 1.
5. Repeat it for the region composed of the points with Z-values 3 and 12.
6. Repeat it for the region composed of the points with Z-values 12 and 15.
7. Repeat it for the region composed of the points with Z-values 1, 3, 9, and 11.
8. Repeat it for the region composed of the points with Z-values 3, 6, 9, and 12.
9. Repeat it for the region composed of the points with Z-values 9, 11, 12, and 14.
10. Repeat it for the region composed of the points with Z-values 8, 9, 10, and 11.

Exercise 28.3 This exercise also refers to Figure 28.3.

1. Consider the region represented by the 01 child of the root in the Region Quad tree shown in Figure 28.3. What are the Z-values of points in this region?
2. Repeat the preceding exercise for the region represented by the 10 child of the root and the 01 child of the 00 child of the root.
3. List the Z-values of four adjacent data points distributed across the four children of the root in the Region Quad tree.
4. Consider the alternative approaches of indexing a two-dimensional point dataset using a B+ tree index: (i) on the composite search key (X, Y) , (ii) on the Z-ordering computed over the X and Y values. Assuming that X and Y values can be represented using two bits each, show an example dataset and query illustrating each of these cases:
 - (a) The alternative of indexing on the composite query is faster.
 - (b) The alternative of indexing on the Z-value is faster.

Exercise 28.4 Consider the Grid file instance with three points 1, 2, and 3 shown in the first part of Figure 28.5.

1. Show the Grid file after inserting each of these points, in the order they are listed: 6, 9, 10, 7, 8, 4, and 5.
2. Assume that deletions are handled by simply removing the deleted points, with no attempt to merge empty or underfull pages. Can you suggest a simple concurrency control scheme for Grid files?
3. Discuss the use of Grid files to handle region data.

Exercise 28.5 Answer each of the following questions independently with respect to the R tree shown in Figure 28.6. (That is, don't consider the insertions corresponding to other questions when answering a given question.)

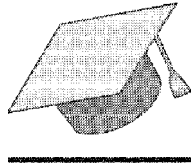
1. Show the bounding box of a new object that can be inserted into R_4 but not into R_3 .
2. Show the bounding box of a new object that is contained in both R_1 and R_6 but is inserted into R_6 .
3. Show the bounding box of a new object that is contained in both R_1 and R_6 and is inserted into R_1 . In which leaf node is this object placed?
4. Show the bounding box of a new object that could be inserted into either R_4 or R_5 but is placed in R_5 based on the principle of least expansion of the bounding box area.

5. Given an example of an object such that searching for the object takes us to both the R1 and R2 subtrees.
6. Give an example query that takes us to nodes R3 and R5. (Explain if there is no such query.)
7. Give an example query that takes us to nodes R3 and R4 but not to R5. (Explain if there is no such query.)
8. Give an example query that takes us to nodes R3 and R5 but not to R4. (Explain if there is no such query.)

BIBLIOGRAPHIC NOTES

Several multidimensional indexing techniques have been proposed. These include B+ files [286], Grid files [565], hB trees [491], KDB trees [630], Pyramid trees [80], Quad trees [649], R trees [350], R* trees [72], R+ trees, the TV tree, and the VA file [767]. [322] discusses how to search R trees for regions defined by linear constraints. Several variations of these, and several other distinct techniques, have also been proposed; Samet's text [650] deals with many of them. A good recent survey is [294].

The use of Hilbert curves for linearizing multidimensional data is proposed in [263]. [118] is an early paper discussing spatial joins. Hellerstein, Naughton, and Pfeffer propose a generalized tree index that can be specialized to obtain many of the specific tree indexes mentioned earlier [376]. Concurrency control and recovery issues for this generalized index are discussed in [447]. Hellerstein, Koutsoupias, and Papadimitriou discuss the complexity of indexing schemes [377], in particular range queries, and Beyer et al. discuss the problems arising with high dimensionality [93]. Faloutsos provides a good overview of how to search multimedia databases by content [258]. A recent trend is towards spatiotemporal applications, such as tracking moving objects [782].



29

FURTHER READING

- ☛ What is next?
- ☛ Key concepts: TP monitors, real-time transactions; data integration; mobile data; main memory databases; multimedia databases; GIS; temporal databases; Bioinformatics; information visualization

This is not the end. It is not even the beginning of the end. But it is, perhaps, the end of the beginning.

-----Winston Churchill

In this book, we concentrated on relational database systems and discussed several fundamental issues in detail. However, our coverage of the database area, and indeed even the relational database area, is far from exhaustive. In this chapter, we look briefly at several topics we did not cover, with the goal of giving the reader some perspective and indicating directions for further study.

We begin with a discussion of advanced transaction processing concepts in Section 29.1. We discuss integrated access to data from multiple databases in Section 29.2 and touch on mobile applications that connect to databases in Section 29.3. We consider the impact of increasingly larger main memory sizes in Section 29.4. We discuss multimedia databases in Section 29.5, geographic information systems in Section 29.6, temporal data in Section 29.7, and sequence data in Section 29.8. We conclude with a look at information visualization in Section 29.9.

The applications covered in this chapter push the limits of currently available database technology and drive the development of new techniques. As even our brief coverage indicates, much work lies ahead for the database field!

29.1 ADVANCED TRANSACTION PROCESSING

The concept of a transaction has wide applicability for a variety of distributed computing tasks, such as airline reservations, inventory management, and electronic Commerce.

29.1.1 Transaction Processing Monitors

Complex applications are often built on top of several resource managers, such as database management systems, operating systems, user interfaces, and messaging software. A transaction processing (TP) monitor glues together the services of several resource managers and provides application programmers a uniform interface for developing transactions with the ACID properties. In addition to providing a uniform interface to the services of different resource managers, a TP monitor also routes transactions to the appropriate resource managers. Finally, a TP monitor ensures that an application behaves as a transaction by implementing concurrency control, logging, and recovery functions and by exploiting the transaction processing capabilities of the underlying resource managers.

TP monitors are used in environments where applications require advanced features, such as access to multiple resource managers, sophisticated request routing (also called workflow management); assigning priorities to transactions and doing priority-based load-balancing across servers, and so on. A DBMS provides many of the functions supported by a TP monitor in addition to processing queries and database updates efficiently. A DBMS is appropriate for environments where the wealth of transaction management capabilities provided by a TP monitor is not necessary and, in particular, where very high scalability (with respect to transaction processing activity) and interoperability are not essential.

The transaction processing capabilities of database systems are improving continually. For example, many vendors offer distributed DBMS products today in which a transaction can execute across several resource managers, each of which is a DBMS. Currently, all the DBMSs must be from the same vendor; however, as transaction-oriented services from different vendors become more standardized, distributed, heterogeneous DBMSs should become available. Eventually, perhaps, the functions of current rrp monitors will also be available in many

DBMSs; for now, TP monitors provide essential infrastructure for high-end transaction processing environments.

29.1.2 New Transaction Models

Consider an application such as computer-aided design, in which users retrieve large design objects from a database and interactively analyze and modify them. Each transaction takes a long time—minutes or even hours, whereas the TPC benchmark transactions take under a millisecond—and holding locks this long affects performance. Further, if a crash occurs, undoing an active transaction completely is unsatisfactory, since considerable user effort may be lost. Ideally, we want to restore most of the actions of an active transaction and resume execution. Finally, if several users are concurrently developing a design, they may want to see changes being made by others without waiting until the end of the transaction that changes the data.

To address the needs of long-duration activities, several refinements of the transaction concept have been proposed. The basic idea is to treat each transaction as a collection of related subtransactions. Subtransactions can acquire locks, and the changes made by a subtransaction become visible to other transactions after the subtransaction ends (and before the main transaction of which it is a part commits). In multilevel transactions, locks held by a subtransaction are released when the subtransaction ends. In nested transactions, locks held by a subtransaction are assigned to the parent (sub)transaction when the subtransaction ends. These refinements to the transaction concept have a significant effect on concurrency control and recovery algorithms.

29.1.3 Real-Time DBMSs

Some transactions must be executed within a user-specified deadline. A hard deadline means the value of the transaction is zero after the deadline. For example, in a DBMS designed to record bets on horse races, a transaction placing a bet is worthless once the race begins. Such a transaction should not be executed; the bet should not be placed. A soft deadline calls the value of the transaction decreases after the deadline, eventually going to zero. For example, in a DBMS designed to monitor some activity (e.g., a complex reactor), a transaction that looks up the current reading of a sensor must be executed within a short time, say, one second. The longer it takes to execute the transaction, the less useful the reading becomes. In a real-time DBMS, the goal is to maximize the value of executed transactions, and the DBMS must prioritize transactions, taking their deadlines into account.

29.2 DATA INTEGRATION

As databases proliferate, users want to access data from more than one source. For example, if several travel agents market their travel packages through the Web, customers would like to compare packages from different agents. A more traditional example is that large organizations typically have several databases, created (and maintained) by different divisions, such as Sales, Production, and Purchasing. While these databases contain much common information, determining the exact relationship between tables in different databases can be a complicated problem. For example, prices in one database might be in dollars per dozen items, while prices in another database might be in dollars per item. The development of XML DTDs (see Section 7.4.3) offers the promise that such *semantic mismatches* can be avoided if all parties conform to a single standard DTD. However, there are many legacy databases and most domains still do not have agreed-upon DTDs; the problem of semantic mismatches will be encountered frequently for the foreseeable future.

Semantic mismatches can be resolved and hidden from users by defining relational views over the tables from the two databases. Defining a collection of views to give a group of users a uniform presentation of relevant data from multiple databases is called *semantic integration*. Creating views that mask semantic mismatches in a natural manner is a difficult task and has been widely studied. In practice, the task is made harder because the schemas of existing databases are often poorly documented; hence, it is difficult to even understand the meaning of rows in existing tables, let alone define unifying views across several tables from different databases.

If the underlying databases are managed using different DBMSs, as is often the case, some kind of ‘middleware’ must be used to evaluate queries over the integrating views, retrieving data at query execution time by using protocols such as Open Database Connectivity (ODBC) to give each underlying database a uniform interface, as discussed in Chapter 6. Alternatively, the integrating views can be materialized and stored in a data warehouse, as discussed in Chapter 25. Queries can then be executed over the warehoused data without accessing the source DBMSs at run-time.

29.3 MOBILE DATABASES

The availability of portable computers and wireless communications has created a new breed of nomadic database users. At one level, these users are simply accessing a database through a network, which is similar to distributed DBMSs. At another level, the network as well as data and user characteristics now have several novel properties, which affect basic assumptions in many components

of a DBMS, including the query engine, transaction manager, and recovery manager:

- Users are connected through a wireless link whose bandwidth is 10 times less than Ethernet and 100 times less than ATM networks. Communication costs are therefore significantly higher in proportion to I/O and CPU costs.
- Users' locations constantly change, and mobile computers have a limited battery life. Therefore, the true communication costs reflect connection time and battery usage in addition to bytes transferred and change constantly depending on location. Data is frequently replicated to minimize the cost of accessing it from different locations.
- As a user moves around, data could be accessed from multiple database servers within a single transaction. The likelihood of losing connections is also much greater than in a traditional network. Centralized transaction management may therefore be impractical, especially if some data is resident at the mobile computers. We may in fact have to give up on ACID transactions and develop alternative notions of consistency for user programs.

29.4 MAIN MEMORY DATABASES

The price of main memory is now low enough that we can buy enough main memory to hold the entire database for many applications; with 64-bit addressing, modern CPUs also have very large address spaces. Some commercial systems now have several *gigabytes* of main memory. This shift prompts a reexamination of some basic DBMS design decisions, since disk accesses no longer dominate processing time for a memory-resident database:

- Main memory does not survive system crashes, and so we still have to implement logging and recovery to ensure transaction atomicity and durability. Log records must be written to stable storage at commit time, and this process could become a bottleneck. To minimize this problem, rather than commit each transaction as it completes, we can collect completed transactions and commit them in batches; this is called group commit. Recovery algorithms can also be optimized, since pages rarely have to be written out to make room for other pages.
- The implementation of in-memory operations has to be optimized carefully, since disk accesses are no longer the limiting factor for performance.
- A new criterion must be considered while optimizing queries, the amount of space required to execute a plan. It is important to minimize the space

overhead because exceeding available physical memory would lead to swapping pages to disk (through the operating system's virtual memory mechanism), greatly slowing down execution.

- Page-oriented data structures become less important (since pages are no longer the unit of data retrieval), and clustering is not important (since the cost of accessing any region of memory is uniform).

29.5 MULTIMEDIA DATABASES

In an object-relational DBMS, users can define ADTs with appropriate methods, which is an improvement over an RDBMS. Nonetheless, supporting just ADTs falls short of what is required to deal with very large collections of multimedia objects, including audio, images, free text, text marked up in HTML or variants, sequence data, and videos. Illustrative applications include NASA's EGS project, which aims to create a repository of satellite imagery; the J. Craig Venter Genome project, which is creating databases of genetic information such as GenBank; and NSF/DARPA's Digital Libraries project, which aims to put entire libraries into database systems and make them accessible through computer networks. Industrial applications, such as collaborative development of engineering designs, also require multimedia database management and are being addressed by several vendors.

We outline some applications and challenges in this area:

- **Content-Based Retrieval:** Users must be able to specify selection conditions based on the contents of multimedia objects. For example, users may search for images using queries such as "Find all images that are similar to this image" and "Find all images that contain at least three airplanes." As images are inserted into the database, the DBMS must analyze them and automatically *extract features* that help answer such content-based queries. This information can then be used to search for images that satisfy a given query, as discussed in Chapter 28. As another example, users would like to search for documents of interest using information retrieval techniques and keyword searches. Vendors are moving toward incorporating such techniques into DBMS products. It is still not clear how these domain-specific retrieval and search techniques can be combined effectively with traditional DBMS queries. Research into abstract data types and ORDBMS query processing has provided a starting point, but more work is needed.
- **Managing Repositories of Large Objects:** Traditionally, DBMSs have concentrated on tables that contain a large number of tuples, each of which is relatively small. Once multimedia objects such as images, sound clips, and videos are stored in a database, individual objects of very large size

have to be handled efficiently. For example, compression techniques must be carefully integrated into the DBMS environment. As another example, distributed DBMSs must develop techniques to efficiently retrieve such objects. Retrieval of multimedia objects in a distributed system has been addressed in limited contexts, such as client-server systems, but in general remains a difficult problem.

- **Video-On-Demand:** Many companies want to provide video-on-demand services that enable users to dial into a server and request a particular video. The video must then be delivered to the user's computer in real time, reliably and inexpensively. Ideally, users must be able to perform familiar VCR functions such as fast-forward and reverse. From a database perspective, the server has to contend with specialized real-time constraints; video delivery rates must be synchronized at the server and at the client, taking into account the characteristics of the communication network.

29.6 GEOGRAPHIC INFORMATION SYSTEMS

Geographic Information Systems (GIS) contain spatial information about cities, states, countries, streets, highways, lakes, rivers, and other geographical features and support applications to combine such spatial information with non-spatial data. As discussed in Chapter 28, spatial data is stored in either raster or vector formats. In addition, there is often a temporal dimension, as when we measure rainfall at several locations over time. An important issue with spatial datasets is how to integrate data from multiple sources, since each source may record data using a different coordinate system to identify locations.

Now let us consider how spatial data in a GIS is analyzed. Spatial information is most naturally thought of as being overlaid on maps. Typical queries include "What cities lie on I-94 between Madison and Chicago?" and "What is the shortest route from Madison to St. Louis?" These kinds of queries can be addressed using the techniques discussed in Chapter 28. An emerging application is in-vehicle navigation aids. With Global Positioning System (GPS) technology, a car's location can be pinpointed, and by accessing a database of local maps, a driver can receive directions from his or her current location to a desired destination; this application also involves mobile database access!

In addition, many applications involve interpolating measurements at certain locations across an entire region to obtain a *model* and combining overlapping models. For example, if we have measured rainfall at certain locations, we can use the Triangulated Irregular Network (TIN) approach to triangulate the region, with the locations at which we have measurements being the vertices of the triangles. Then, we use some form of interpolation to estimate

the rainfall at points within triangles. Interpolation, triangulation, map overlays, visualization of spatial data, and many other domain-specific operations are supported in GIS products such as ESRI Systems' ARC-Info. Therefore, while spatial query processing techniques as discussed in Chapter 28 are an important part of a GIS product, considerable additional functionality must be incorporated as well. How best to extend RDBMS systems with this additional functionality is an important problem yet to be resolved. Agreeing on standards for data representation formats and coordinate systems is another major challenge facing the field.

29.7 TEMPORAL DATABASES

Consider the following query: "Find the longest interval in which the same person managed two different departments." Many issues are associated with representing temporal data and supporting such queries. We need to be able to distinguish the times during which something is true in the real world (**valid time**) from the times it is true in the database (**transaction time**). The period during which a given person managed a department can be indicated by two fields *from* and *to*, and queries must reason about time intervals. Further, temporal queries require the DBMS to be aware of the anomalies associated with calendars (such as leap years).

29.8 BIOLOGICAL DATABASES

Bioinformatics is an emerging field at the intersection of Biology and Computer Science. From a database standpoint, the rapidly growing data in this area has (at least) two interesting characteristics. First, a lot of *loosely structured data* is widely exchanged, leading to interest in integration of such data. This has motivated some of the research in the area of XML repositories.

The second interesting feature is *sequence data*. DNA sequences are being generated at a rapid pace by the biological community. The field of biological information management and analysis has become very popular in recent years, called bioinformatics. Biological data, such as DNA sequence data, characterized by complex structure and numerous relationships among data elements, many overlapping and incomplete or erroneous data fragments (because experimentally collected data from several groups, often working on related problems, is stored in the databases), a need to frequently change the database *schema* itself as new kinds of relationships in the data are discovered, and the need to maintain several versions of data for archival and reference.

29.9 INFORMATION VISUALIZATION

As computers became faster and main memory cheaper, it becomes increasingly feasible to create visual presentations of data, rather than just text-based reports. Data visualization makes it easier for users to understand the information in large complex datasets. The challenge here is to make it easy for users to develop visual presentations of their data and interactively query such presentations. Although a number of data visualization tools are available, efficient visualization of large datasets presents many challenges.

The need for visualization is especially important in the context of decision support; when confronted with large quantities of high-dimensional data and various kinds of data summaries produced by using analysis tools such as SQL, OLAP, and data mining algorithms, the information can be overwhelming. Visualizing the data, together with the generated summaries, can be a powerful way to sift through this information and spot interesting trends or patterns. The human eye, after all, is very good at finding patterns. A good framework for data mining must combine analytic tools to process data and bring out latent anomalies or trends with a visualization environment in which a user can notice these patterns and interactively drill down to the original data for further analysis.

29.10 SUMMARY

The database area continues to grow vigorously, in terms of both technology and applications. The fundamental reason for this growth is that the amount of information stored and processed using computers is growing rapidly. Regardless of the nature of the data and the intended applications, users need database management systems and their services (concurrent access, crash recovery, easy and efficient querying, etc.) as the volume of data increases. As the range of applications is broadened, however, some shortcomings of current DBMSs become serious limitations. These problems are being actively studied in the database research community.

The coverage in this book provides an introduction, but is not intended to cover all aspects of database systems. Ample material is available for further study, as this chapter illustrates, and we hope that the reader is motivated to pursue the leads in the bibliography. Bon voyage!

BIBLIOGRAPHIC NOTES

[338] contains a comprehensive treatment of all aspects of transaction processing. See [241] for several papers that describe new transaction models for nontraditional applications such as CAD/CAM. [1,577,696,711,761] are some of the many papers on real-time databases.

Determining which entities are the same across different databases is a difficult problem; it is an example of a semantic mismatch. Resolving such mismatches has been addressed in many papers, including [424, 476, 641, 663]. [389] is an overview of theoretical work in this area. Also see the bibliographic notes for Chapter 22 for references to related work on multidatabases, and see the notes for Chapter 2 for references to work on view integration.

[304] is an early paper on main memory databases. [102, 406] describe the Dali main memory storage manager. [421] surveys visualization idioms designed for large databases, and [342] discusses visualization for data mining.

Visualization systems for databases include DataSpace [592], DEVise [489], IVEE [27], the Mineset suite from SGI, Tioga [31], and VisDB [420]. In addition, a number of general tools are available for data visualization.

Querying text repositories has been studied extensively in information retrieval; see [626] for a recent survey. This topic has generated considerable interest in the database community recently because of the widespread use of the Web, which contains many text sources. In particular, HTML documents have some structure if we interpret links as edges in a graph. Such documents are examples of semistructured data; see [2] for a good overview. Recent papers on queries over the Web include [2, 445, 527, 564].

See [576] for a survey of multimedia issues in database management. There has been much recent interest in database issues in a mobile computing environment; for example, [387,398]. See [395] for a collection of articles on this subject. [728] contains several articles that cover all aspects of temporal databases. The use of constraints in databases has been actively investigated in recent years; [416] is a good overview. Geographic Information Systems have also been studied extensively; [586] describes the Paradise system, which is notable for its scalability.

The book [794] contains detailed discussions of temporal databases (including the TSQL2 language, which is influencing the SQL standard), spatial and multimedia databases, and uncertainty in databases.

THE MINIBASE SOFTWARE

Practice is the best of all instructors.

-Publius Syrus, 42 B.C.

Minibase is a small relational DBMS, together with a suite of visualization tools, that has been developed for use with this book. While the book makes no direct reference to the software and can be used independently, Minibase offers instructors an opportunity to design a variety of hands-on assignments, with or without programming. To see an online description of the software, visit this URL:

<http://www.cs.wisc.edu/-dbbook/minibase.html>

The software is available freely through ftp. By registering themselves as users at the URL for the book, instructors can receive prompt notification of any major bug reports and fixes. Sample project assignments, which elaborate on some of the briefly sketched ideas in the *project-based exercises* at the end of chapters, can be seen at

<http://www.cs.wisc.edu/-dbbook/minihwk.html>

Instructors should consider making all modifications to each assignment to discourage undesirable 'code reuse' by students; assignment handouts formatted using Latex are available by ftp. Instructors can also obtain solutions to these assignments by contacting the authors (raghu@cs.wisc.edu, johannes@cs.cornell.edu).

30.1 WHAT IS AVAILABLE

Minibase is intended to supplement the use of a commercial DBMS such as Oracle or Sybase in course projects, not to replace them. While a commercial DBMS is ideal for SQL assignments, it does not help students understand how the DBMS works. Minibase is intended to address the latter issue; the subset of SQL that it supports is intentionally kept small, and students should also be asked to use a commercial DBMS for writing SQL queries and programs.

Minibase is provided on an as-is basis with no warranties or restrictions for educational or personal use. It includes the following:

- Code for a small single-user relational DBMS, including a parser and query optimizer for a subset of SQL, and components designed to be (re)written by students as project assignments: *heap files*, *buffer manager*, *B+ trees*, *sorting*, and *joins*.

30.2 OVERVIEW OF MINIBASE ASSIGNMENTS

Several assignments involving the use of Minibase are described here. Each of these has been tested in a course already, but the details of how Minibase is set up might vary at your school, so you may have to modify the assignments accordingly. If you plan to use these assignments, you are advised to download and try them at your site well in advance of handing them to students. We have done our best to test and document these assignments and the Minibase software, but bugs undoubtedly persist. Please report bugs at this URL:

<http://www.cs.wisc.edu/-dbbook/minibase.comments.html>

We hope users will contribute bug fixes, additional project assignments, and extensions to Minibase. These will be made publicly available through the Minibase site, together with pointers to the authors.

In several assignments, students are asked to rewrite a component of Minibase. The book provides the necessary background for all these assignments, and the assignment handout provides additional system-level details. The online HTML documentation provides an overview of the software, in particular the component interfaces, and can be downloaded and installed at each school that uses Minibase. The projects that follow should be assigned after covering the relevant material from the indicated chapter:

- **Buffer Manager (Chapter 9):** Students are given code for the layer that manages space on disk and supports the concept of pages with page ids. They are asked to implement a buffer manager that brings requested pages into memory if they are not already there. One variation of this assignment could use different replacement policies. Students are asked to assemble a single-user environment, with no concurrency control or recovery management.
- **HF Page (Chapter 9):** Students must write code that manages records on a page using a slot-directory page format to keep track of the records. Possible variants include fixed-length versus variable-length records and other ways to keep track of records on a page.

- **Heap Files (Chapter 9):** Using the HF page and buffer manager code, students are asked to implement a layer that supports the abstraction of files of unordered pages, that is, heap files.
- **B+ Trees (Chapter 10):** This is one of the more complex assignments. Students have to implement a page class that maintains records in sorted order within a page and implement the B+ tree index structure to impose a sort order across several leaf-level pages. Indexes store $\langle key, record\text{-}pointer \rangle$ pairs in leaf pages, and data records are stored separately (in heap files). Similar assignments can easily be created for Linear Hashing or Extendible Hashing index structures.
- **External sorting (Chapter 13):** Building on the buffer manager and heap file layers, students are asked to implement external merge-sort. The emphasis is on minimizing I/O rather than on the in-memory sort used to create sorted runs.
- **Sort-Merge Join (Chapter 14):** Building upon the code for external sorting, students are asked to implement the sort-merge join algorithm. This assignment can be easily modified to create assignments that involve other join algorithms.
- **Index Nested-Loop Join (Chapter 14):** This assignment is similar to the sort-merge join assignment, but relies on B+ tree (or other indexing) code, instead of sorting code.

30.3 ACKNOWLEDGMENTS

The Minibase software was inspired by Minirel, a small relational DBMS developed by David DeWitt for instructional use. Minibase was developed by a large number of dedicated students over a long time, and the design was guided by Mike Carey and R. Ramakrishnan. See the online documentation for more on Minibase's history.

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