

### **Chapter 15: Query Processing**

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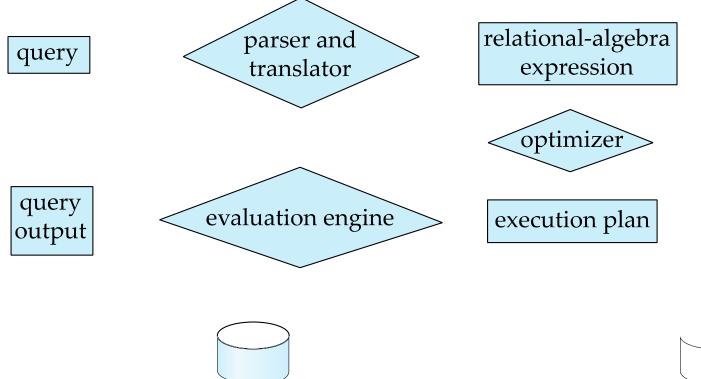
### **Chapter 15: Query Processing**

- Overview
- Measures of Query Cost
- Selection Operation
- Sorting
- Join Operation
- Other Operations
- Evaluation of Expressions



## **Basic Steps in Query Processing**

- 1. Parsing and translation
- 2. Optimization
- 3. Evaluation



# Basic Steps in Query Processing (Cont.)

- Parsing and translation
  - translate the query into its internal form. This is then translated into relational algebra.
  - Parser checks syntax, verifies relations
- Evaluation
  - The query-execution engine takes a query-evaluation plan, executes that plan, and returns the answers to the query.



### Basic Steps in Query Processing: Optimization

- A relational algebra expression may have many equivalent expressions
  - E.g.,  $\sigma_{salary < 75000}(\prod_{salary}(instructor))$  is equivalent to  $\prod_{salary}(\sigma_{salary < 75000}(instructor))$
- Each relational algebra operation can be evaluated using one of several different algorithms
  - Correspondingly, a relational-algebra expression can be evaluated in many ways.
- Annotated expression specifying detailed evaluation strategy is called an evaluation-plan. E.g.:
  - Use an index on *salary* to find instructors with salary < 75000,
  - Or perform complete relation scan and discard instructors with salary ≥ 75000



## **Basic Steps: Optimization (Cont.)**

- Query Optimization: Amongst all equivalent evaluation plans choose the one with lowest cost.
  - Cost is estimated using statistical information from the database catalog
    - e.g. number of tuples in each relation, size of tuples, etc.
- In this chapter we study
  - How to measure query costs
  - Algorithms for evaluating relational algebra operations
  - How to combine algorithms for individual operations in order to evaluate a complete expression
- In Chapter 16
  - We study how to optimize queries, that is, how to find an evaluation plan with lowest estimated cost



## **Measures of Query Cost**

- Many factors contribute to time cost
  - disk access, CPU, and network communication
- Cost can be measured based on
  - **response time**, i.e. total elapsed time for answering query, or
  - total resource consumption
- We use total resource consumption as cost metric
  - Response time harder to estimate, and minimizing resource consumption is a good idea in a shared database
- We ignore CPU costs for simplicity
  - Real systems do take CPU cost into account
  - Network costs must be considered for parallel systems
- We describe how estimate the cost of each operation
  - We do not include cost to writing output to disk



### **Measures of Query Cost**

- Disk cost can be estimated as:
  - Number of seeks \* average-seek-cost
  - Number of blocks read \* average-block-read-cost
  - Number of blocks written \* average-block-write-cost
- For simplicity we just use the number of block transfers from disk and the number of seeks as the cost measures
  - $t_{T}$  time to transfer one block
    - Assuming for simplicity that write cost is same as read cost
  - $t_{\rm S}$  time for one seek
  - Cost for b block transfers plus S seeks
     b \* t<sub>T</sub> + S \* t<sub>S</sub>
- $t_{s}$  and  $t_{\tau}$  depend on where data is stored; with 4 KB blocks:
  - High end magnetic disk:  $t_s = 4$  msec and  $t_T = 0.1$  msec
  - SSD: :  $t_{S}$  = 20-90 microsec and  $t_{T}$  = 2-10 microsec for 4KB



## Measures of Query Cost (Cont.)

- Required data may be buffer resident already, avoiding disk I/O
  - But hard to take into account for cost estimation
- Several algorithms can reduce disk IO by using extra buffer space
  - Amount of real memory available to buffer depends on other concurrent queries and OS processes, known only during execution
- Worst case estimates assume that no data is initially in buffer and only the minimum amount of memory needed for the operation is available
  - But more optimistic estimates are used in practice



## **Selection Operation**

#### File scan

- Algorithm A1 (linear search). Scan each file block and test all records to see whether they satisfy the selection condition.
  - Cost estimate =  $b_r$  block transfers + 1 seek
    - *b<sub>r</sub>* denotes number of blocks containing records from relation *r*
  - If selection is on a key attribute, can stop on finding record
    - cost =  $(b_r/2)$  block transfers + 1 seek
  - Linear search can be applied regardless of
    - selection condition or
    - ordering of records in the file, or
    - availability of indices
- Note: binary search generally does not make sense since data is not stored consecutively
  - except when there is an index available,
  - and binary search requires more seeks than index search



## **Selections Using Indices**

- Index scan search algorithms that use an index
  - selection condition must be on search-key of index.
- A2 (clustering index, equality on key). Retrieve a single record that satisfies the corresponding equality condition
  - $Cost = (h_i + 1) * (t_T + t_S)$
- A3 (clustering index, equality on nonkey) Retrieve multiple records.
  - Records will be on consecutive blocks
    - Let b = number of blocks containing matching records

• 
$$Cost = h_i * (t_T + t_S) + t_S + t_T * b$$



## **Selections Using Indices**

- A4 (secondary index, equality on key/non-key).
  - Retrieve a single record if the search-key is a candidate key
    - $Cost = (h_i + 1) * (t_T + t_S)$
  - Retrieve multiple records if search-key is not a candidate key
    - each of n matching records may be on a different block
    - Cost =  $(h_i + n) * (t_T + t_S)$ 
      - Can be very expensive!



## **Selections Involving Comparisons**

- Can implement selections of the form  $\sigma_{A \le V}(r)$  or  $\sigma_{A \ge V}(r)$  by using
  - a linear file scan,
  - or by using indices in the following ways:
- A5 (clustering index, comparison). (Relation is sorted on A)
  - For  $\sigma_{A \ge V}(r)$  use index to find first tuple  $\ge v$  and scan relation sequentially from there
  - For σ<sub>A≤V</sub>(r) just scan relation sequentially till first tuple > v; do not use index
- A6 (clustering index, comparison).
  - For  $\sigma_{A \ge V}(r)$  use index to find first index entry  $\ge v$  and scan index sequentially from there, to find pointers to records.
  - For σ<sub>A≤V</sub>(r) just scan leaf pages of index finding pointers to records, till first entry > v
  - In either case, retrieve records that are pointed to
  - requires an I/O per record; Linear file scan may be cheaper!



## **Implementation of Complex Selections**

- Conjunction:  $\sigma_{\theta 1} \wedge \theta_{\theta 2} \wedge \dots \theta_{\theta n}(r)$
- A7 (conjunctive selection using one index).
  - Select a combination of  $\theta_i$  and algorithms A1 through A7 that results in the least cost for  $\sigma_{\theta i}$  (*r*).
  - Test other conditions on tuple after fetching it into memory buffer.
- A8 (conjunctive selection using composite index).
  - Use appropriate composite (multiple-key) index if available.
- A9 (conjunctive selection by intersection of identifiers).
  - Requires indices with record pointers.
  - Use corresponding index for each condition, and take intersection of all the obtained sets of record pointers.
  - Then fetch records from file
  - If some conditions do not have appropriate indices, apply test in memory.

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## **Algorithms for Complex Selections**

- Disjunction:  $\sigma_{\theta 1} \vee_{\theta 2} \vee \ldots \otimes_{\theta n} (r)$ .
- A10 (disjunctive selection by union of identifiers).
  - Applicable if *all* conditions have available indices.
    - Otherwise use linear scan.
  - Use corresponding index for each condition, and take union of all the obtained sets of record pointers.
  - Then fetch records from file
- Negation: σ<sub>¬θ</sub>(r)
  - Use linear scan on file
  - If very few records satisfy  $\neg \theta$ , and an index is applicable to  $\theta$ 
    - Find satisfying records using index and fetch from file



## **Bitmap Index Scan**

- The **bitmap index scan** algorithm of PostgreSQL
  - Bridges gap between secondary index scan and linear file scan when number of matching records is not known before execution
  - Bitmap with 1 bit per page in relation
  - Steps:
    - Index scan used to find record ids, and set bit of corresponding page in bitmap
    - Linear file scan fetching only pages with bit set to 1
  - Performance
    - Similar to index scan when only a few bits are set
    - Similar to linear file scan when most bits are set
    - Never behaves very badly compared to best alternative



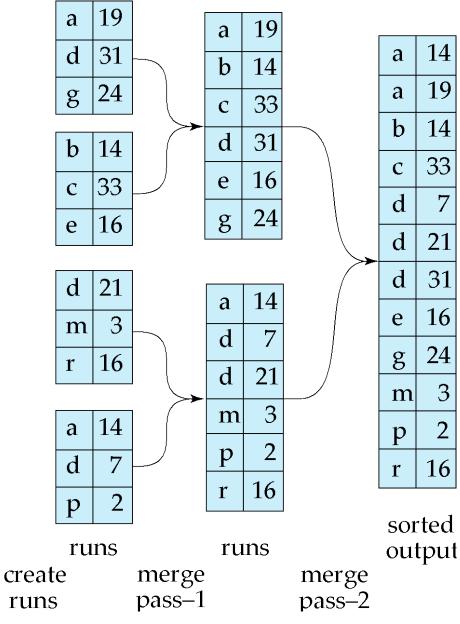
## Sorting

- We may build an index on the relation, and then use the index to read the relation in sorted order. May lead to one disk block access for each tuple.
- For relations that fit in memory, techniques like quicksort can be used.
   For relations that don't fit in memory, external sort-merge is a good choice.



### **Example: External Sorting Using Sort-Merge**

24 g 19 a d 31 33 С b 14 16 e 16 r d 21 3 m 2 p d 7 14 а initial relation



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### **External Sort-Merge**

Let *M* denote memory size (in pages).

1. Create sorted runs. Let *i* be 0 initially.

Repeatedly do the following till the end of the relation:

- (a) Read *M* blocks of relation into memory
- (b) Sort the in-memory blocks
- (c) Write sorted data to run  $R_i$ ; increment *i*.

Let the final value of *i* be *N* 

2. Merge the runs (next slide).....



## **External Sort-Merge (Cont.)**

#### 2. Merge the runs (N-way merge). We assume (for now) that N < M.

1. Use *N* blocks of memory to buffer input runs, and 1 block to buffer output. Read the first block of each run into its buffer page

#### 2. repeat

- 1. Select the first record (in sort order) among all buffer pages
- 2. Write the record to the output buffer. If the output buffer is full write it to disk.
- Delete the record from its input buffer page.
   If the buffer page becomes empty then read the next block (if any) of the run into the buffer.
- **3. until** all input buffer pages are empty:



## **External Sort-Merge (Cont.)**

- If  $N \ge M$ , several merge *passes* are required.
  - In each pass, contiguous groups of *M* 1 runs are merged.
  - A pass reduces the number of runs by a factor of M -1, and creates runs longer by the same factor.
    - E.g. If M=11, and there are 90 runs, one pass reduces the number of runs to 9, each 10 times the size of the initial runs
  - Repeated passes are performed till all runs have been merged into one.



## **External Merge Sort (Cont.)**

- Cost analysis:
  - 1 block per run leads to too many seeks during merge
    - Instead use b<sub>b</sub> buffer blocks per run
      - → read/write  $b_b$  blocks at a time
    - Can merge  $\lfloor M/b_b \rfloor$  –1 runs in one pass
  - Total number of merge passes required:  $\lceil \log_{\lfloor M/bb \rfloor 1}(b_r/M) \rceil$ .
  - Block transfers for initial run creation as well as in each pass is  $2b_r$ 
    - for final pass, we don't count write cost
      - we ignore final write cost for all operations since the output of an operation may be sent to the parent operation without being written to disk
    - Thus total number of block transfers for external sorting:  $b_r (2 \lceil \log_{\lfloor M/bb \rfloor - 1} (b_r / M) \rceil + 1) \rceil$
  - Seeks: next slide



## **External Merge Sort (Cont.)**

- Cost of seeks
  - During run generation: one seek to read each run and one seek to write each run
    - 2[b<sub>r</sub>/M]
  - During the merge phase
    - Need  $2 \lceil b_r / b_b \rceil$  seeks for each merge pass
      - except the final one which does not require a write
    - Total number of seeks:

 $2 \lceil b_r / M \rceil + \lceil b_r / b_b \rceil (2 \lceil \log_{\lfloor M/bb \rfloor - 1} (b_r / M) \rceil - 1)$ 



## **Join Operation**

- Several different algorithms to implement joins
  - Nested-loop join
  - Block nested-loop join
  - Indexed nested-loop join
  - Merge-join
  - Hash-join
- Choice based on cost estimate
- Examples use the following information
  - Number of records of *student*: 5,000 *takes*: 10,000
  - Number of blocks of *student*: 100 *takes*: 400



### **Nested-Loop Join**

- To compute the theta join r ⋈ θ s
   for each tuple t<sub>r</sub> in r do begin
   for each tuple t<sub>s</sub> in s do begin
   test pair (t<sub>r</sub>, t<sub>s</sub>) to see if they satisfy the join condition θ
   if they do, add t<sub>r</sub> t<sub>s</sub> to the result.
   end
   end
- *r* is called the **outer relation** and *s* the **inner relation** of the join.
- Requires no indices and can be used with any kind of join condition.
- Expensive since it examines every pair of tuples in the two relations.



## **Nested-Loop Join (Cont.)**

In the worst case, if there is enough memory only to hold one block of each relation, the estimated cost is

 $n_r * b_s + b_r$  block transfers, plus  $n_r + b_r$  seeks

- If the smaller relation fits entirely in memory, use that as the inner relation.
  - Reduces cost to  $b_r + b_s$  block transfers and 2 seeks
- Assuming worst case memory availability cost estimate is
  - with *student* as outer relation:
    - 5000 \* 400 + 100 = 2,000,100 block transfers,
    - 5000 + 100 = 5100 seeks
  - with *takes* as the outer relation
    - 10000 \* 100 + 400 = 1,000,400 block transfers and 10,400 seeks
- If smaller relation (*student*) fits entirely in memory, the cost estimate will be 500 block transfers.
- Block nested-loops algorithm (next slide) is preferable.

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## **Block Nested-Loop Join**

 Variant of nested-loop join in which every block of inner relation is paired with every block of outer relation.

```
for each block B_r of r do begin
for each block B_s of s do begin
for each tuple t_r in B_r do begin
for each tuple t_s in B_s do begin
Check if (t_r, t_s) satisfy the join condition
if they do, add t_r \cdot t_s to the result.
end
end
end
```



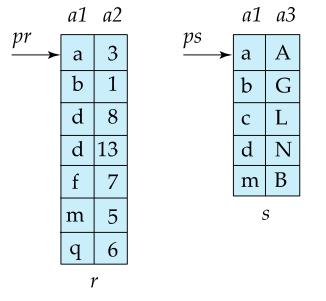
### **Indexed Nested-Loop Join**

- Index lookups can replace file scans if
  - join is an equi-join or natural join and
  - an index is available on the inner relation's join attribute
    - Can construct an index just to compute a join.
- For each tuple t<sub>r</sub> in the outer relation r, use the index to look up tuples in s that satisfy the join condition with tuple t<sub>r</sub>.
- Worst case: buffer has space for only one page of *r*, and, for each tuple in *r*, we perform an index lookup on *s*.
- Cost of the join:  $b_r (t_T + t_S) + n_r * c$ 
  - Where c is the cost of traversing index and fetching all matching s tuples for one tuple or r
  - c can be estimated as cost of a single selection on s using the join condition.
- If indices are available on join attributes of both r and s, use the relation with fewer tuples as the outer relation.



## **Merge-Join**

- 1. Sort both relations on their join attribute (if not already sorted on the join attributes).
- 2. Merge the sorted relations to join them
  - 1. Join step is similar to the merge stage of the sort-merge algorithm.
  - 2. Main difference is handling of duplicate values in join attribute every pair with same value on join attribute must be matched
  - 3. Detailed algorithm in book





## Merge-Join (Cont.)

- Can be used only for equi-joins and natural joins
- Each block needs to be read only once (assuming all tuples for any given value of the join attributes fit in memory
- Thus the cost of merge join is:
  - $b_r + b_s$  block transfers  $+ \lfloor b_r / b_b \rfloor + \lfloor b_s / b_b \rfloor$  seeks
  - + the cost of sorting if relations are unsorted.
- hybrid merge-join: If one relation is sorted, and the other has a secondary B<sup>+</sup>-tree index on the join attribute
  - Merge the sorted relation with the leaf entries of the B+-tree .
  - Sort the result on the addresses of the unsorted relation's tuples
  - Scan the unsorted relation in physical address order and merge with previous result, to replace addresses by the actual tuples
    - Sequential scan more efficient than random lookup

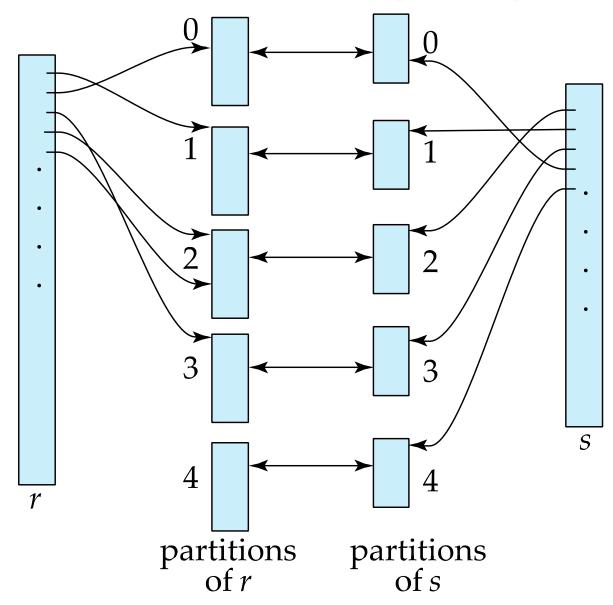


### Hash-Join

- Applicable for equi-joins and natural joins.
- A hash function *h* is used to partition tuples of both relations
- h maps JoinAttrs values to {0, 1, ..., n}, where JoinAttrs denotes the common attributes of r and s used in the natural join.
  - $r_0, r_1, \ldots, r_n$  denote partitions of r tuples
    - Each tuple  $t_r \in r$  is put in partition  $r_i$  where  $i = h(t_r [JoinAttrs])$ .
  - *r*<sub>0</sub>, *r*<sub>1</sub>..., *r*<sub>n</sub> denotes partitions of *s* tuples
    - Each tuple  $t_s \in s$  is put in partition  $s_i$ , where  $i = h(t_s [JoinAttrs])$ .
- Note: In book, Figure 12.10 r<sub>i</sub> is denoted as H<sub>ri</sub>, s<sub>i</sub> is denoted as H<sub>si</sub> and
   n is denoted as n<sub>h</sub>



### Hash-Join (Cont.)





## **Hash-Join Algorithm**

The hash-join of *r* and *s* is computed as follows.

- 1. Partition the relation *s* using hashing function *h*. When partitioning a relation, one block of memory is reserved as the output buffer for each partition.
- 2. Partition *r* similarly.
- 3. For each *i*:
  - (a) Load  $s_i$  into memory and build an in-memory hash index on it using the join attribute. This hash index uses a different hash function than the earlier one *h*.
  - (b) Read the tuples in  $r_i$  from the disk one by one. For each tuple  $t_r$  locate each matching tuple  $t_s$  in  $s_i$  using the inmemory hash index. Output the concatenation of their attributes.

Relation *s* is called the **build input** and *r* is called the **probe input**.



## Hash-Join algorithm (Cont.)

- The value n and the hash function h is chosen such that each s<sub>i</sub> should fit in memory.
  - Typically n is chosen as [b<sub>s</sub>/M] \* f where f is a "fudge factor", typically around 1.2
  - The probe relation partitions  $s_i$  need not fit in memory
- Recursive partitioning required if number of partitions n is greater than number of pages M of memory.
  - instead of partitioning n ways, use M-1 partitions for s
  - Further partition the M 1 partitions using a different hash function
  - Use same partitioning method on *r*
  - Rarely required: e.g., with block size of 4 KB, recursive partitioning not needed for relations of < 1GB with memory size of 2MB, or relations of < 36 GB with memory of 12 MB</li>



## **Handling of Overflows**

- Partitioning is said to be skewed if some partitions have significantly more tuples than some others
- Hash-table overflow occurs in partition s<sub>i</sub> if s<sub>i</sub> does not fit in memory.
   Reasons could be
  - Many tuples in s with same value for join attributes
  - Bad hash function
- Overflow resolution can be done in build phase
  - Partition  $s_i$  is further partitioned using different hash function.
  - Partition  $r_i$  must be similarly partitioned.
- Overflow avoidance performs partitioning carefully to avoid overflows during build phase
  - E.g. partition build relation into many partitions, then combine them
- Both approaches fail with large numbers of duplicates
  - Fallback option: use block nested loops join on overflowed partitions



## **Cost of Hash-Join**

- If recursive partitioning is not required: cost of hash join is  $3(b_r + b_s) + 4 * n_h$  block transfers +  $2(\lceil b_r/b_b \rceil + \lceil b_s/b_b \rceil)$  seeks
- If recursive partitioning required:
  - number of passes required for partitioning build relation *s* to less than M blocks per partition is  $\lceil log_{\lfloor M/bb \rfloor 1}(b_s/M) \rceil$
  - best to choose the smaller relation as the build relation.
  - Total cost estimate is:  $2(b_r + b_s) \lceil log_{\lfloor M/bb \rfloor - 1}(b_s/M) \rceil + b_r + b_s \text{ block transfers } + 2(\lceil b_r/b_b \rceil + \lceil b_s/b_b \rceil) \lceil log_{\lfloor M/bb \rfloor - 1}(b_s/M) \rceil \text{ seeks}$
- If the entire build input can be kept in main memory no partitioning is required
  - Cost estimate goes down to  $b_r + b_s$ .



# **Hybrid Hash–Join**

- Useful when memory sized are relatively large, and the build input is bigger than memory.
- Main feature of hybrid hash join: Keep the first partition of the build relation in memory.
- E.g. With memory size of 25 blocks, *instructor* can be partitioned into five partitions, each of size 20 blocks.
  - Division of memory:
    - The first partition occupies 20 blocks of memory
    - 1 block is used for input, and 1 block each for buffering the other 4 partitions.
- teaches is similarly partitioned into five partitions each of size 80
  - the first is used right away for probing, instead of being written out
- Cost of 3(80 + 320) + 20 +80 = 1300 block transfers for hybrid hash join, instead of 1500 with plain hash-join.
- Hybrid hash-join most useful if  $M >> \sqrt{b_s}$



# **Complex Joins**

Join with a conjunctive condition:

 $r \bowtie_{\theta 1 \land \theta 2 \land \dots \land \theta n} s$ 

- Either use nested loops/block nested loops, or
- Compute the result of one of the simpler joins  $r \bowtie_{\theta i} s$ 
  - final result comprises those tuples in the intermediate result that satisfy the remaining conditions

$$\theta_1 \wedge \ldots \wedge \theta_{i-1} \wedge \theta_{i+1} \wedge \ldots \wedge \theta_n$$

Join with a disjunctive condition

$$r \bowtie_{\theta_1 \lor \theta_2 \lor \ldots \lor \theta_n} s$$

- Either use nested loops/block nested loops, or
- Compute as the union of the records in individual joins  $r \bowtie_{\theta_i} s$ :  $(r \bowtie_{\theta_1} s) \cup (r \bowtie_{\theta_2} s) \cup \ldots \cup (r \bowtie_{\theta_n} s)$

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# **Joins over Spatial Data**

- No simple sort order for spatial joins
- Indexed nested loops join with spatial indices
  - R-trees, quad-trees, k-d-B-trees



# **Other Operations**

- Duplicate elimination can be implemented via hashing or sorting.
  - On sorting duplicates will come adjacent to each other, and all but one set of duplicates can be deleted.
  - *Optimization:* duplicates can be deleted during run generation as well as at intermediate merge steps in external sort-merge.
  - Hashing is similar duplicates will come into the same bucket.

## Projection:

- perform projection on each tuple
- followed by duplicate elimination.



# **Other Operations : Aggregation**

- Aggregation can be implemented in a manner similar to duplicate elimination.
  - **Sorting** or **hashing** can be used to bring tuples in the same group together, and then the aggregate functions can be applied on each group.
  - Optimization: partial aggregation
    - combine tuples in the same group during run generation and intermediate merges, by computing partial aggregate values
    - For count, min, max, sum: keep aggregate values on tuples found so far in the group.
      - When combining partial aggregate for count, add up the partial aggregates
    - For avg, keep sum and count, and divide sum by count at the end



# **Other Operations : Set Operations**

- Set operations (∪, ∩ and —): can either use variant of merge-join after sorting, or variant of hash-join.
- E.g., Set operations using hashing:
  - 1. Partition both relations using the same hash function
  - 2. Process each partition *i* as follows.
    - 1. Using a different hashing function, build an in-memory hash index on  $r_i$ .
    - 2. Process s<sub>i</sub> as follows
      - $r \cup s$ :
        - 1. Add tuples in  $s_i$  to the hash index if they are not already in it.
        - 2. At end of  $s_i$  add the tuples in the hash index to the result.



# **Other Operations : Set Operations**

- E.g., Set operations using hashing:
  - 1. as before partition *r* and *s*,
  - 2. as before, process each partition *i* as follows
    - 1. build a hash index on  $r_i$
    - 2. Process s<sub>i</sub> as follows
      - *r* ∩ s:
        - 1. output tuples in  $s_i$  to the result if they are already there in the hash index
      - r s:
        - 1. for each tuple in  $s_{i}$ , if it is there in the hash index, delete it from the index.
        - 2. At end of  $s_i$  add remaining tuples in the hash index to the result.



# **Answering Keyword Queries**

- Indices mapping keywords to documents
  - For each keyword, store sorted list of document IDs that contain the keyword
    - Commonly referred to as a inverted index
    - E.g.: database: d1, d4, d11, d45, d77, d123 distributed: d4, d8, d11, d56, d77, d121, d333
  - To answer a query with several keywords, compute intersection of lists corresponding to those keywords
- To support ranking, inverted lists store extra information
  - **"Term frequency**" of the keyword in the document
  - "Inverse document frequency" of the keyword
  - Page rank of the document/web page



# **Other Operations : Outer Join**

- Outer join can be computed either as
  - A join followed by addition of null-padded non-participating tuples.
  - by modifying the join algorithms.
- Modifying merge join to compute  $r \bowtie s$ 
  - In  $r \bowtie s$ , non participating tuples are those in  $r \prod_R (r \bowtie s)$
  - Modify merge-join to compute  $r \bowtie s$ :
    - During merging, for every tuple t<sub>r</sub> from r that do not match any tuple in s, output t<sub>r</sub> padded with nulls.
  - Right outer-join and full outer-join can be computed similarly.



# **Other Operations : Outer Join**

- Modifying hash join to compute  $r \bowtie s$ 
  - If r is probe relation, output non-matching r tuples padded with nulls
  - If *r* is build relation, when probing keep track of which *r* tuples matched *s* tuples. At end of *s<sub>i</sub>* output non-matched *r* tuples padded with nulls



# **Evaluation of Expressions**

- So far: we have seen algorithms for individual operations
- Alternatives for evaluating an entire expression tree
  - Materialization: generate results of an expression whose inputs are relations or are already computed, materialize (store) it on disk. Repeat.
  - Pipelining: pass on tuples to parent operations even as an operation is being executed
- We study above alternatives in more detail

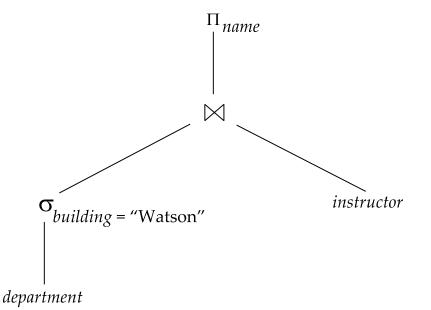


## **Materialization**

- Materialized evaluation: evaluate one operation at a time, starting at the lowest-level. Use intermediate results materialized into temporary relations to evaluate next-level operations.
- E.g., in figure below, compute and store

 $\sigma_{building="Watson"}(department)$ 

then compute the store its join with *instructor*, and finally compute the projection on *name*.



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# Materialization (Cont.)

- Materialized evaluation is always applicable
- Cost of writing results to disk and reading them back can be quite high
  - Our cost formulas for operations ignore cost of writing results to disk, so
    - Overall cost = Sum of costs of individual operations + cost of writing intermediate results to disk
- Double buffering: use two output buffers for each operation, when one is full write it to disk while the other is getting filled
  - Allows overlap of disk writes with computation and reduces execution time



# Pipelining

- Pipelined evaluation: evaluate several operations simultaneously, passing the results of one operation on to the next.
- E.g., in previous expression tree, don't store result of

 $\sigma_{building="Watson"}(department)$ 

- instead, pass tuples directly to the join.. Similarly, don't store result of join, pass tuples directly to projection.
- Much cheaper than materialization: no need to store a temporary relation to disk.
- Pipelining may not always be possible e.g., sort, hash-join.
- For pipelining to be effective, use evaluation algorithms that generate output tuples even as tuples are received for inputs to the operation.
- Pipelines can be executed in two ways: demand driven and producer driven



# Pipelining (Cont.)

- In demand driven or lazy evaluation
  - system repeatedly requests next tuple from top level operation
  - Each operation requests next tuple from children operations as required, in order to output its next tuple
  - In between calls, operation has to maintain "state" so it knows what to return next

### In producer-driven or eager pipelining

- Operators produce tuples eagerly and pass them up to their parents
  - Buffer maintained between operators, child puts tuples in buffer, parent removes tuples from buffer
  - if buffer is full, child waits till there is space in the buffer, and then generates more tuples
- System schedules operations that have space in output buffer and can process more input tuples
- Alternative name: **pull** and **push** models of pipelining



# Pipelining (Cont.)

- Implementation of demand-driven pipelining
  - Each operation is implemented as an iterator implementing the following operations
    - open()
      - E.g. file scan: initialize file scan
        - state: pointer to beginning of file
      - E.g.merge join: sort relations;
        - state: pointers to beginning of sorted relations
    - next()
      - E.g. for file scan: Output next tuple, and advance and store file pointer
      - E.g. for merge join: continue with merge from earlier state till

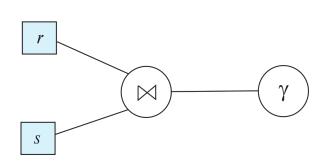
next output tuple is found. Save pointers as iterator state.

close()

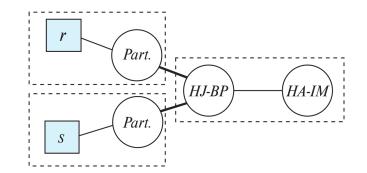


# **Blocking Operations**

- Blocking operations: cannot generate any output until all input is consumed
  - E.g. sorting, aggregation, ...
- But can often consume inputs from a pipeline, or produce outputs to a pipeline
- Key idea: blocking operations often have two suboperations
  - E.g. for sort: run generation and merge
  - For hash join: partitioning and build-probe
- Treat them as separate operations







(b) Pipelined Plan

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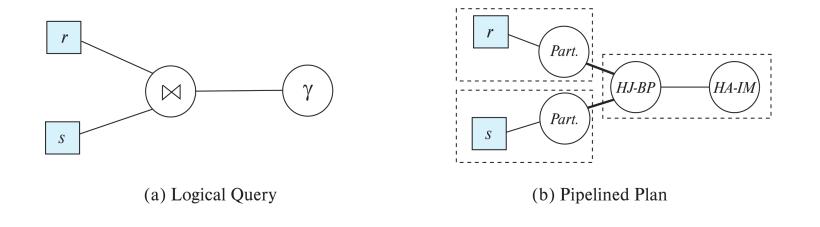
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# **Pipeline Stages**

### Pipeline stages:

- All operations in a stage run concurrently
- A stage can start only after preceding stages have completed execution



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# **Evaluation Algorithms for Pipelining**

- Some algorithms are not able to output results even as they get input tuples
  - E.g. merge join, or hash join
  - intermediate results written to disk and then read back
- Algorithm variants to generate (at least some) results on the fly, as input tuples are read in
  - E.g. hybrid hash join generates output tuples even as probe relation tuples in the in-memory partition (partition 0) are read in
  - **Double-pipelined join technique**: Hybrid hash join, modified to buffer partition 0 tuples of both relations in-memory, reading them as they become available, and output results of any matches between partition 0 tuples
    - When a new r<sub>0</sub> tuple is found, match it with existing s<sub>0</sub> tuples, output matches, and save it in r<sub>0</sub>
    - Symmetrically for s<sub>0</sub> tuples



# **Pipeling for Continuous-Stream Data**

### Data streams

- Data entering database in a continuous manner
- E.g. Sensor networks, user clicks, ...

### Continuous queries

- Results get updated as streaming data enters the database
- Aggregation on windows is often used
  - E.g. **tumbling windows** divide time into units, e.g. hours, minutes
- Need to use pipelined processing algorithms
  - **Punctuations** used to infer when all data for a window has been received



# **Query Processing in Memory**

- Query compilation to machine code
  - Overheads of interpretation
    - E.g. repeatedly finding attribute location within tuple, from metadata
    - Overhead of expression evaluation
  - Compilation can avoid many such overheads and speed up query processing
  - Often via generation of Java byte code / LLVM, with just-in-time (JIT) compilation
- Column-oriented storage
  - Allows vector operations (in conjunction with compilation)
- Cache conscious algorithms



# **Cache Conscious Algorithms**

- Goal: minimize cache misses, make best use of data fetched into the cache as part of a cache line
- For sorting:
  - Use runs that are as large as L3 cache (a few megabytes) to avoid cache misses during sorting of a run
  - Then merge runs as usual in merge-sort
- For hash-join
  - First create partitions such that build+probe partitions fit in memory
  - Then subpartition further s.t. build subpartition+index fits in L3 cache
    - Speeds up probe phase significantly by avoiding cache misses
- Lay out attributes of tuples to maximize cache usage
  - Attributes that are often accessed together should be stored adjacent to each other
- Use multiple threads for parallel query processing
  - Cache misss leads to stall of one thread, but others can proceed

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## **End of Chapter 15**