### CMSC 424 – Database design Lecture 17 Query processing

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# Admin

- Homework 3 is on the website
- Project part 1 due
- Midterm answers/results
   > 90 A (6)
   75-90 B (9)
   50-75 C (6)
   < 50 D (1)</li>
- 10 improved, 10 got worse, 2 about the same

# Wake up...skills you should have

- Find information online (e.g. what is this BibTex after all)
- Write a parser for a simple file format
- Adapt/incorporate an existing parser
- Manage your time (start early, evaluate difficulty of project)
- Pay attention in class
- Communicate

## **Complex Selections**

- conjunction  $\sigma_{\theta_1 \land \theta_2}$  s1 = # of tuples satisfying  $\theta_1$ s2 = # of tuples satisfying  $\theta_2$ combined SC = s1 \* s2/(n(r) \* n(r))assuming independence of predicates
- disjunction  $\sigma_{\theta_1 \lor \theta_2}$

combined SC = 
$$1 - (1 - s1/n(r)) * (1 - s2/n(r))$$

- this 1 minus the probability of all predicates are satisfied at once (s1/n(r) + s2/n(r) s1/n(r)\*s2/n(r))- union of results
- negation  $\sigma_{\neg_{\theta}}$

$$n(\sigma_{\neg \theta}(r)) = n(r) - n(\sigma_{\theta}(r))$$

# Multiple Index Selection

<u>GOAL</u>: apply the most restrictive one and combine multiple of them to reduce the intermediate results AS EARLY AS POSSIBLE

- conjunctive selection using one index A: select using A and then apply the remaining of the predicates on the retrieved tuple values
- conjunctive selection using a composite key index (R.A,R.B)- then create a composite key or range from the query values and search directly (range search on the first attribute only)
- conjunctive selection using two indexes A and B: search each separately and intersect the tuple identifiers (TIDs)
- disjunctive selection using two indexes A and B: search each separately and take the union of the TIDs

### Join Methods: Nested Loop

- tuple-oriented:

   for each tuple t(r) in r do begin
   for each tuple t(s) in s do begin
   join(t(r),t(s) and append the result to the output
   end
- block-oriented:

   for each block b(r) in r do begin
   for each block b(s) in s do begin
   join(b(r),b(s) and append the result to the output
   end
   end
   inner Î
- reverse inner loop similar to above but for even outer blocks we scan the inner relation in reverse



outer blocks



#### Cost of Block-Oriented Nested Loop Buffer size M+1

- cost depends on the number of buffers and the buffer replacement strategy
  - fasten 1 block from the outer relation, M for the inner and LRU

cost: b(r) + b(r)\*b(s) assuming that b(s) > M

- fasten M blocks from the outer relation, and 1 for the inner
- 1: read M from the outer cost: M blocks
- 2: for each block of s join 1 X M blocks cost: b(s) -"-
- 3: repeat with the next M blocks of r until all done repeated b(r)/M times

 $cost = [ (M + b(s)]^* b(r)/M = b(r)+[b(r)^*b(s)]/M$ 

• which relation should be the outer?

### Join Methods: Sort-Merge-Join

- two phases
  - sorting phase: sort both relations (this can be done in parallel)
  - merging phase: join tuples during the merge

sort R on joining attribute sort S on joining attribute merge(sorted-R,sorted-S)

• cost with M buffers

$$\operatorname{cost} = \underbrace{b_r \left(2 \left\lceil \log_{M-1}(b_r/M) \right\rceil + 1\right)}_{\operatorname{Sorting R}} + \underbrace{b_r + b_s}_{K} \left(2 \left\lceil \log_{M-1}(b_s/M) \right\rceil + 1\right) + \underbrace{b_s}_{Write} + \underbrace{b_r + b_s}_{Write} \xrightarrow{Write}_{Merge}$$

if one pass is required the expressions  $\lceil \log M - 1(br/M) \rceil = 1$  and  $\lceil \log M - 1(bs/M) \rceil = 1$  so the total cost is 3\*b(r)+b(r)+b(r)+3\*b(s)+b(s)+b(s) = 5\*b(r)+5\*b(s)However, if M> b(r) and b(s) then the expression evaluates to 3\*b(r)+3\*b(s)).

# Join Methods: Hash-Join

- two phases
  - hash phase: hash both relations into hashed partitions (this can be done in parallel)
  - bucket-wise join phase: join tuples of the same partitions only

hash R on the joining into H(R) buckets hash S on the joining into H(S) buckets nested-loop join of corresponding buckets Hj(R),Hj(S) or main-memory hash index join of -"-

 Number of partitions is large to make each partition of Hj(R) fit in the buffer memory
 -- each Hj(R) consists of several blocks



 We assume that buckets of Hj(R) fit in the buffer memory (and that after hashing the partitions of R and S have the same size with R and S):

cost = b(r) + b(r) + b(s) + b(s) + b(r) + b(s) = 3 (b(r) + b(s))

#### Hash-Join Algorithm Details 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 inmory hash index. Output the concatenation of their attributes.

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

# Example of Cost of Hash-Join

- M= 20 blocks
- $b_{depositor} = 100$

 $customer \bowtie depositor$ 

- $b_{customer} = 400.$
- *depositor* is the build input. Partition it into 5 partitions, each of size 20 blocks. This partitioning can be done in one pass.
- partition *customer* into 5 partitions, each of size 80. This is also done in one pass.
- Do the partition joins- for each j put 20 blocks of partition depositor(j) in memory, built the hash index, and do the probes with the 80 blocks of customer(j)
- Therefore total cost, ignoring cost of writing partially filled blocks:

-3(100 + 400) = 1500 I/Os

# 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  $r_i$  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., recursive partitioning not needed for relations of 1GB or less with memory size of 2MB, with block size of 4KB.

### Join Methods: Indexed-Join

• inner relation has an index (clustering or not)

```
for each block b(r) in r do begin
for each tuple t(r) in b(r) do begin
search the index B on s with the value t.A of the joining attr. A
and join(t(r),t.A)
end
end
```

```
    cost = b(r) + n(r)* cost(σ(S.A=c))
where cost(σ(S.A=c)) is as computed for indexed selection
```

### Estimation of Join Size: "(R 🖂 S)

•  $0 \le n(\mathbf{R} \bowtie \mathbf{S}) \le n(r)^* n(s)$  (0 when nothing joins and when everything joins)

- if joining attribute is a key of R then n(R ⋈S) <= n(s) /\* each value of S.A would join to at most one value of R.A \*/

- if -"- is a key of R and a foreign key of S then  $n(R \bowtie S) = n(s)$ /\* each value of S.A would join to exactly one value of R.A \*/

- if is not a key then each value of A in R appears n(s)/V(A,s) times in S, therefore, n(r) tuples of R produce:  $n(R \bowtie S) = n(r)*n(s) / V(A,s)$ symmetrically we can obtain:  $n(R \bowtie S) = n(r)*n(s) / V(A,r)$ 

if the values are different we use:  $\min\{n(r)^*n(s) / V(A,s), n(r)^*n(s) / V(A,r)\}$ 

# **Other Operations**

- Outer Joins
  - Left outer join easy
  - Right/Full outer join (may need some bookkeeping)
- Duplicate elimination
  - Hard
  - Sort at the end and eliminate
  - Hash output and eliminate
- Aggregates
  - Sum, count, min, max easily kept during execution
  - -Avg = Sum / count
  - Std = sqrt(ssum/count)

# **External Sorting with Sort-Merge**

•external vs internal sorting: relation/file does not fit in memory

•create runs phase: repeat until done

read M blocks of the relation (or rest if <=M)

internal sort using any sort method, e.g QuickSort(M)

write the sorted tuples into a run R data file

end

•merge-runs phase:

read one block from each run;

merge tuples on the result;

advance the pointer from the run you appended last;

- if the block of a run is empty, read the next one until all blocks of all runs are done
  - this assumes that a block from each run can be kept in main memory. If not, then the same algorithm has to be applied in multiple passes



# **External Merge Sort Cost**

- Cost analysis:
  - Initial number of runs:  $b_r/M$
  - Total number of merge passes required:  $\lceil \log_{M-1}(b_r/M) \rceil$ .
  - Block transfers for initial run creation is  $b_r + b_r = 2b_r$ 
    - 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 pipelined to the display or to a parent operation without being written to disk. If pipelined, it will be counted in the cost of the follow up operator

• Thus total number of block transfers for external sorting:

 $2 b_r (\lceil \log_{M-1}(b_r/M) \rceil) + b_r = b_r (2 \lceil \log_{M-1}(b_r/M) \rceil + 1)$ 

- If  $M \ge \lceil b_r/M \rceil$  (only one pass is required) the expression  $\lceil \log_{M-1}(b_r/M) \rceil = 1$ total cost =  $3b_r$
- However, if  $M > b_r$  then this expression evaluates to 0 total cost = $b_r$  ONLY