

Data Storage

- What is stored: Schemas, Relation metadata (e.g. indexes, statistical info), Log files

Secondary Storage

- DBMS storage includes:
 - Data: Stored in disk blocks/pages
 - File layer: Organisation and data retrieval
 - Buffer manager: Reading/writing of disk pages
 - Disk space manager: Keeps track of pages used by file layer

Magnetic Hard-Disk Drive (HDD)

- Disk Access time:
 - Command Processing time (negligible)
 - Seek time: Move disk head on track
 - Rotational delay: Rotate to put head on start of correct sector
 - Avg rotational delay = time for $\frac{1}{2}$ revolution = $\frac{1}{2} \times \frac{60}{\text{RPM}}$ s
 - Transfer time: Rotate along the correct sector(s) to move data to/from disk = $\frac{\text{\# sectors read}}{\text{Total \# sectors on track}} \times \frac{60}{\text{RPM}}$ s
- Sequential I/O:
 - Pages stored contiguously on one track, then move on to next surface of the cylinder (i.e. same track across different surfaces), then move on to next cylinder
- Solid-State Drive (SSD)**
 - Per block: Avg seek time + Avg rotational delay + Transfer time

Buffer Manager

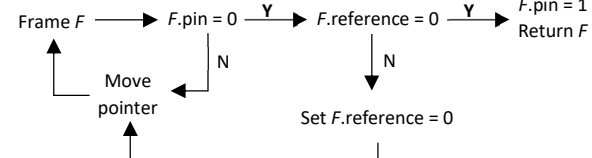
- Maintains a buffer pool in RAM (memory allocated for DBMS) for caching. Each unit of memory is called a **frame**
 - Each frame has:
 - Pin count: # clients using the page
 - Dirty flag: Whether page has been modified but not updated on disk
- Disk pages are fetched into/release from the buffer pool
- Page request procedure:
 - Client requests page P
 - Is page P already in memory?
 - Yes: Pin frame F . Return address of F . **End**
 - No: continue to 3
 - Find free frame or evict a page if buffer pool is full (i.e. Find some frame F . **pin = 0**)
 - Pin frame F
 - Write frame F into disk if F is dirty
 - Return address of frame F

Page Replacement Policies

Clock Replacement Policy

- Each frame has an additional reference bit
- Pointer moves in FIFO manner
- Only frame with reference bit = 0 **AND** pin = 0 will be replaced
- Reference bit set to 0 when pin drops to 0

- Clock replacement procedure:

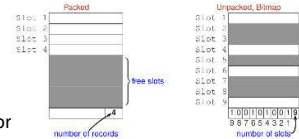


- DO NOT** increment pointer after each successful page replacement

Page/Record Format

Fixed-length Records

- Each record is of fixed length.
- Each record is identified by a RID = (page id, slot #)
- Packed: Contiguous;
- Unpacked: Not Contiguous
- Record Format: Fields in a record



Variable-length Records (Slotted Page Organisation)

- Slot directory:
 - Pointer to start of free space: Memory location for next record to be inserted
 - Number of slots: Increments whenever a new record is inserted
 - Pointers to each record: Address to the start of a record in the page + size of the record (byte offset)
- Record Format:
 - Delimit fields with special symbols (e.g. F1 \$ F2 \$... \$ Fn)
 - Use an array of field offsets

Tree-based Indexing

- Index:** Data structure to speed up data retrieval; stored as a file.)
- Search key:** Sequence of k data attributes e.g. (name, age)
- Unique index:** Search key is a candidate key
- Data entry:** (Key Value, RID), stored in index

B+ Tree Index

- Leaf nodes** store data entries. Leaf nodes are doubly-linked and all of them are on the same level
- Height h :** # levels of internal nodes (root is level 0). Leaf nodes are at level h
- Internal nodes** store index entries in the form $(p_0, k_1, p_1, k_2, p_2, \dots, p_n)$
- Order** of a B+ Tree, d : Root node must contain $[1, 2d]$ entries and non-root nodes must contain $[d, 2d]$ entries
- A B+ Tree with n level of internal nodes has with order d has $2(d+1)^{n-1} \leq \text{\# leafs} \leq (2d+1)^n$

Search

- At each internal node, find largest k_i s.t. target $k_i \leq k$.
 - Search subtree at p_i if k_i exists
 - Otherwise, search subtree at p_0
- Continue until leaf node and return all entries with search key = k .
 - If range search, traverse along leaf nodes and return all entries within the bound

Insert (Handling overflows)

- Node splitting**
 - Overflowing leaf node**
 - Distribute the $d+1$ largest entries into new leaf node
 - Create and insert new index entry using smallest key in new leaf node into parent node
 - If parent node overflows, split parent node
 - Overflowing internal node**
 - Split at the middle key, and push it up to the parent node
 - Propagate node splitting until no overflows/reached root
- Redistribution of data entries**
 - Redistribute entries in overflowed leaf node N by putting the largest /smallest entry (among the $2d+1$ entries) into adjacent right/left sibling N'
 - Then, update the separating key in parent

Delete (Handling underflows)

- Node Merging**
 - Underflowing leaf node**
 - Merge underflowed leaf node N with adjacent sibling N' by moving all entries from N' to N .
 - Then, delete N' and the separating key in parent
 - Update parent index if needed
 - Underflowing internal node** (Pre-condition: N' must have d entries)
 - Merge underflowed internal node N with adjacent sibling N' by pulling down separating key in parent, combining N and N'
 - Propagate node merging until no underflows/reached root
- Redistribution of data entries** (Pre-condition: N' must have $> d$ entries)
 - Redistribute entries by moving the data entry with the smallest/largest key from right/left sibling N' to underflowing leaf node N
 - Update separating key in parent with the smallest key in N'
- Redistribution of internal entries**
 - Merge leaf nodes, causing internal node N to be underflowed
 - "Pull down" the separating parent key K between N and sibling N' and join with N
 - Then, replace the K in the affected index entry in parent node with $N'.k_i$ (i.e. left/right-most key in right/left sibling)
 - Remove k_i from sibling

Data Formats

- Format-1:** Leaves store data records
- Format-2:** Leaves store (k, rid)
- Format-3:** Leaves store (k, rid-list)

Bulk Loading

- Sort data entries to be inserted by search key
 - Load the leaf pages with those sorted entries
 - Initialize the B+ tree with an empty root page
 - For each leaf page (in sequential order), insert its index entry into the rightmost parent-of-leaf level page of the B+ tree
- Advantages:
 - Efficient construction
 - Leaf pages are allocated sequentially

Hash-based Indexing

- Numbers in the buckets are the **HASH VALUES**, NOT THE KEY VALUES

Linear Hashing

- N_0 = initial # of buckets = 2^m
- N_i = # buckets in at start of level $i = 2^i N_0 = 2^{m+i}$
- Split image** of $B_j = B_{j+N_i}$

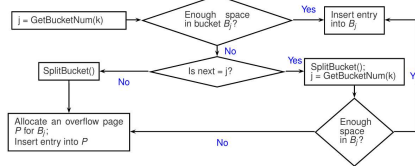
Insert

- Hash function: $h_i(k) = h(k) \bmod N_i$ (look at the **last $m + i$ bits**)

$$\text{Bucket \#} = \begin{cases} h_i(k), & h_i(k) \geq \text{next} \\ h_{i+1}(k), & \text{otherwise} \end{cases}$$

- Splitting:** occurs when any bucket overflows:

- Split the bucket B_j pointed to by 'next'
- Redistribute entries:
 - Entries in B_j : $(m + i + 1)^{\text{th}}$ bit is 0
 - Entries in B_{j+N_i} : $(m + i + 1)^{\text{th}}$ bit is 1



Delete

- If last bucket B_{N_i+next} is empty:
 - If next > 0
 - Decrement next
 - Delete last bucket
 - If next = 0 and level > 0
 - Decrement level
 - Update next to point to last bucket in previous level ($B_{N_{i-1}-1}$)
- Delete overflow pages that become empty after redistribution

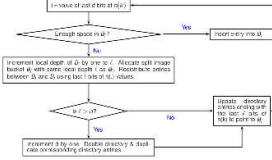
Performance

- Best case: No overflow pages – 1 disk I/O per insertion
- Worst case: All hashed to the same bucket – linear I/O cost
- Average: 1.2 disk I/O per insertion

Extendible Hashing

- Global depth (directory) = d ; # directory entries = 2^d
- Local depth (bucket) = $l \leq d$
- Directory entry # = **last d bits** of $h(k)$; points to bucket
- All entries in a bucket have same l bits in their $h(k)$
- Corresponding entries: differ only by the d^{th} bit (indexed 1)
- # directory entries pointing to a bucket = 2^{d-l}
- Splitting:** occurs when target bucket B_i overflows
 - Increment l of B_i
 - Allocate new bucket B_j (split image) with same l
 - Redistribute:
 - Entries in B_i : l^{th} bit is 0
 - Entries in B_j : l^{th} bit is 1
 - Using the last l bits, redistribute pointer(s) between B_i and B_j

- Bucket B_i overflows:
 - If $l = d$
 - Increment d and l of B_i
 - Double the number of directory entries
 - Split B_i ; redistribute
 - Redistribute pointer(s)
 - If $l \leq d$
 - Increment l of B_i
 - Split B_i ; redistribute
 - Redistribute pointer(s)



Delete

- If entries in B_i and B_j (corresponding entry) can fit into 1 bucket:
 - Merge B_i and B_j into one bucket e.g. B_i
 - Delete B_j
 - Decrement l of B_i
 - Move directory entries pointing to B_j to point to B_i
 - If each pair of corresponding entries point to the same bucket, decrement d and halve the directory

Performance

- At most 2 disk I/O per insertion

Index Formats

Clustered vs Unclustered

- Clustered: Order of data records is close to order of the data entries
 - Format-1 index is a clustered index
 - Cost of RID lookups become 0 (format-1) or $\|\sigma(R)\| / b_d$
- Each relation can have at most one clustered index

Dense vs Sparse

- Dense: There is an index record for every search key value; otherwise, it is sparse
- Unclustered index must be dense
- Format-1 B+ Tree index is sparse

Query Evaluation: Sort

External Mergesort

- Assume N data records need to be sorted
- Assume that data records are on the disk
- Only B memory pages are allocated for sorting
 - 1 page is allocated for **output**
 - $B - 1$ pages are allocated for **input**
- Read the N data records into $[N/B]$ initial sorted runs
- Recursively merge the sorted runs in a $B - 1$ way merge
 - With $B - 1$ input pages, we can sort $B - 1$ sorted runs by allocating one page for each of the $B - 1$ sorted runs
 - Each input pages has a pointer to the current smallest value
 - When output page is full, write the page back to disk

Analysis

- N_0 = # sorted runs in initial pass (pass 0) = $\lceil N/B \rceil$
- Total # passes = $\lceil \log_{B-1}(N_0) \rceil + 1$
- Total # I/O = $2N(\lceil \log_{B-1}(N_0) \rceil + 1)$
 - Each pass has N reads and writes

External Mergesort with Blocked I/O

- Instead of reading/writing one page at a time, read/write in units of **buffer blocks** of b_i and b_o pages respectively
 - This allows for **sequential** I/O over random I/O
- Assume B memory pages are allocated for sorting
 - Output** block size = b_o pages; **Input** block size = b_i pages
 - 1 block is allocated for **output**; $\lceil \frac{B-b_o}{b_i} \rceil$ blocks are allocated for **input**
- Read the N data records into $[N/B]$ sorted runs
- Recursively merge the sorted runs in a $\lceil \frac{B-b_o}{b_i} \rceil$ way merge
 - With $\lceil \frac{B-b_o}{b_i} \rceil$ input blocks, we can sort $\lceil \frac{B-b_o}{b_i} \rceil$ sorted runs by allocating one block for each of the sorted runs
 - When output block is full, write the block back to disk

Analysis

- N_0 = # sorted runs in initial pass (pass 0) = $\lceil N/B \rceil$
- F = # sorted runs that can be merged at each merge pass = $\lceil \frac{B-b_o}{b_i} \rceil$
- Total # passes = $\lceil \log_F(N_0) \rceil + 1$
- Reduced merge factor, but more sequential I/O

Query Evaluation: Select σ

- Access paths:
 - Table scan: Scan all data pages
 - Index scan: Scan index pages + RID lookup (if needed)
 - Index intersection: Combine results from multiple index scans + RID lookup (if needed)
- More selective access path \rightarrow fewer pages need to be accessed
- Include Columns:** Specify the attributes whose values are also stored in the data entries of the index, on top of the index key. Can be used to avoid RID lookups
- Covering Index:** Index I is a covering index for query Q if all attributes in Q are part of the key or include columns(s) of I
 - Then, Q is evaluated with index-only plan
- Matching Predicates**
 - B+ Tree:** $I = (K_1, K_2, \dots, K_n)$. I matches predicate p if p is in the form: $(K_1 = c_1) \wedge \dots \wedge (K_{i-1} = c_{i-1}) \wedge (K_i \text{ op}_i c_i), \quad i \in [1, n]$
 - Prefix key with all equality operator, and at most one non-equality operator on last attribute of prefix
 - Hash Index:** $I = (K_1, K_2, \dots, K_n)$. I matches predicate p if p is in the form: $(K_1 = c_1) \wedge (K_2 = c_2) \wedge \dots \wedge (K_n = c_n)$
 - All attributes must appear in p and only equality operator

Primary Conjuncts p'

- The subset of conjuncts in p that I matches

Covered Conjuncts p_c

- The subset of conjuncts p_c in p such that all attributes in p_c appear in the key or include column(s) of I

Cost Evaluation

B+ Tree Index

$$Cost_{internal} = \begin{cases} \left\lceil \log_F \left\lceil \frac{\|R\|}{b_d} \right\rceil \right\rceil, & \text{format} - 1 \\ \left\lceil \log_F \left\lceil \frac{\|R\|}{b_i} \right\rceil \right\rceil, & \text{otherwise} \end{cases}$$

$$Cost_{leaf} = \begin{cases} \left\lceil \frac{\|\sigma_{pr}(R)\|}{b_d} \right\rceil, & \text{format} - 1 \\ \left\lceil \frac{\|\sigma_{pr}(R)\|}{b_i} \right\rceil, & \text{otherwise} \end{cases}$$

$$Cost_{RID} = \begin{cases} 0, & \text{format} - 1 \text{ or covering} \\ \|\sigma_{pc}(R)\|, & \text{otherwise (worst case)} \\ \frac{\|\sigma_{pc}(R)\|}{b_d}, & \text{otherwise (clustered)} \end{cases}$$

$$Cost_{total} = Cost_{internal} + Cost_{leaf} + Cost_{RID}$$

Hash Index

- Ranged query: Table scan = $|R|$
- Format 1
 - $Cost_{total} = Cost_{records} = \left\lceil \frac{\|\sigma_{pr}(R)\|}{b_d} \right\rceil$
- Format 2
 - $Cost_{entries} \geq \left\lceil \frac{\|\sigma_{pr}(R)\|}{b_i} \right\rceil$, due to possible long overflow chain
 - $Cost_{records} = \begin{cases} 0, & I \text{ is covering index} \\ \|\sigma_{pc}(R)\|, & \text{otherwise} \end{cases}$
 - $Cost_{total} = Cost_{records} + Cost_{entries}$

Query Evaluation: Project π

- $\pi_L(R)$ – No duplicates (select DISTINCT) ; $\pi_L^*(R)$ – Keep duplicates

Sort-Based Projection

Unoptimized Approach

- Extract attributes L from records $R \rightarrow \pi_L^*(R)$
- Sort $\pi_L^*(R)$ using L as sort key \rightarrow sorted $\pi_L^*(R)$ [External Mergesort]
- Scan $\pi_L^*(R)$ to remove duplicates $\rightarrow \pi_L(R)$

Analysis

- Read I/O (table scan) = $|R|$; Write I/O = $|\pi_L^*(R)| \rightarrow |R| + |\pi_L^*(R)|$
- Given B buffer pages for sorting:
 - Merge factor = $B - 1$; $N_0 = |\pi_L^*(R)/B|$
 - Step 2 total = $2|\pi_L^*(R)|(log_{B-1} N_0 + 1)$
- Step 3 total = $|\pi_L^*(R)|$ (ignore write I/O as cost)
- Total = $|R| + 2|\pi_L^*(R)|(log_{B-1} N_0 + 2)$

Optimized Approach

- Create sorted runs with attributes L
 - Write only attributes L to output page
- Merge sorted runs and remove duplicates simultaneously
 - Analysis**
 - Read I/O = $|R|$; Write I/O = $|\pi_L^*(R)| \rightarrow$ Step 1 total = $|R| + |\pi_L^*(R)|$
 - Given B buffer pages for sorting:
 - Merge factor = $B - 1$; $N_0 = |\pi_L^*(R)/B|$
 - Step 2 total = $2|\pi_L^*(R)|(log_{B-1} N_0) - |\pi_L^*(R)|$
 - Total = $|R| + 2|\pi_L^*(R)|(log_{B-1} N_0)$

- If $B > \sqrt{|\pi_L^*(R)|}$,
 - # initial sorted runs = $\left\lceil \frac{|R|}{B} \right\rceil \approx \sqrt{|\pi_L^*(R)|}$
 - # merging passes = $log_{B-1}(\sqrt{|\pi_L^*(R)|}) \approx 1$
 - Total = $|R| + 2|\pi_L^*(R)|$

Hash-Based Projection

- Partition** all the tuples in R into R_1, R_2, \dots, R_{B-1}
 - Only B memory pages are allocated for hash table
 - 1 page is allocated for **input** buffer
 - $B - 1$ pages for **output** buffers/partitions R_1, R_2, \dots, R_{B-1}
 - For each t in R ,
 - $h(\pi_L(t))$, then output $\pi_L(t)$ into output buffer $R_{h(\pi_L(t))}$
 - If $R_{h(\pi_L(t))}$ is full, flush to disk.
 - Output of partition phase is $\pi_L^*(R_1), \pi_L^*(R_2), \dots, \pi_L^*(R_{B-1})$
- Duplicate Elimination:** For each $\pi_L^*(R_i)$,
 - Initialize hash table T of size $B - 1$
 - For each tuple t in $\pi_L^*(R_i)$:
 - 2.2.1 Perform $h'(t) = j$ (NOTE: $h' \neq h$)
 - 2.2.2 If t not in B_j , then insert t into B_j
 - Output all the tuples in T as $\pi_L(R_i)$
- Combine all $\pi_L(R_i)$
- Entire T must fit in main memory. If not, recursively partition $\pi_L^*(R_i)$ into $\pi_L^*(R_{i_1}), \dots, \pi_L^*(R_{i_{B-1}})$

Analysis

- Assuming no partition overflow:
 - Partitioning = $|R| + |\pi_L^*(R)|$
 - Read I/O (table scan) = $|R|$.
 - Each t is projected before writing. So, write I/O = $|\pi_L^*(R)|$.
 - Duplicate elimination = $|\pi_L^*(R)|$
 - Each of the projected tuples is read once. Ignore write I/O
 - Total = $|R| + 2|\pi_L^*(R)|$
- If $B > \sqrt{f|\pi_L^*(R)|}$, then there will not be partition overflow
 - Assume h hashes every t in R uniformly
 - Each R_i will have $\approx \frac{|\pi_L^*(R)|}{B-1}$ pages $\rightarrow B > f \frac{|\pi_L^*(R)|}{B-1} \approx \sqrt{f|\pi_L^*(R)|}$

Indexing

- If there is a covering index I for the projection, then replace table scan (i.e. read I/O = $|R|$) in the both projection schemes with index scan
- If index is ordered (e.g. B+-tree) whose search key (e.g. (A, B, C)) includes wanted attributes (e.g. $\pi_{A,B}(R)$) as a prefix
 - Scan all the data entries in order
 - Compare adjacent data entries for duplicates

Query Evaluation: Join \bowtie

- In general, the smaller relation should be the outer relation

Tuple-based Nested Loop Join

- For each tuple r in R , for each tuple s in S , check if r matches s

Analysis

- Scan $R = |R|$; Scan $S = \|R\| \times |S|$
- Total = $|R| + \|R\| \times |S|$

Page-based Nested Loop Join

- For each page P_r in R , for each page P_s in S , for each tuple r in P_r , for each tuple s in P_s , check if r matches s

Analysis

- Scan $R = |R|$; Scan $S = |R| \times |S|$
- Total = $|R| + |R| \times |S|$

Block Nested Loop Join

- Allocate 1 buffer page for output, 1 buffer page for S and $B - 2$ buffer pages for R . Read pages from R in blocks of $B - 2$
- For each block B_i from R , for each page P_s in S , for each tuple r in B_i , for each tuple s in P_s , check if r matches s

Analysis

- Scan $R = |R|$; Scan $S = \left\lceil \frac{|R|}{B-2} \right\rceil \times |S|$
- Total = $|R| + \left\lceil \frac{|R|}{B-2} \right\rceil \times |S|$

Index Nested Loop Join

- Precondition: There is an index on the join attribute(s) of **inner** relation
- For each tuple r in R , use r to probe S 's index to find matching tuples

Analysis

- Assuming uniform distribution, one tuple in R matches with $\left\lceil \frac{\|S\|}{\|\pi_{B(S)}\|} \right\rceil$ tuples
- Scan $R = |R|$; Scan $S = \|R\| \times \left(\log_F \left\lceil \frac{\|S\|}{b_d} \right\rceil + \left\lceil \frac{\|S\|}{b_d \|\pi_{B(S)}\|} \right\rceil \right)$
- Total = $|R| + \|R\| \times \left(\log_F \left\lceil \frac{\|S\|}{b_d} \right\rceil + \left\lceil \frac{\|S\|}{b_d \|\pi_{B(S)}\|} \right\rceil \right)$

Sort-Merge Join

- Assume $|R| \leq |S|$; R is outer relation

Unoptimized Approach

- Sort:** Sort both R and S based on \bowtie attributes
 - This partitions R into R_1, \dots, R_k and S into S_1, \dots, S_l each containing tuples with the same join attribute value(s)
- Merge:**
 - Initialize pointers p_r and p_s for R and S , each pointing to the first tuple. Let r and s be the tuples pointed to by p_r and p_s
 - r and s do not match: Advance smaller pointer
 - 2.2.1 If p_r is advanced and r matches s at p'_s , **rewind**: $p_s \leftarrow p'_s$
 - r and s match:
 - 2.3.1 Remember position $p_s : p'_s \leftarrow p_s$
 - 2.3.2 Output $r \bowtie s$ and advance p_s until r and s do not match

Analysis

- Sort R and S :
 - External Mergesort = $2|R|(\log_m(N_R) + 1) + 2|S|(\log_m(N_S) + 1)$
 - Internal Mergesort = $|R| + |S|$
- Merge:
 - Best case: No rewinds = $|R| + |S|$
 - Worst case: All tuples in R and S match = $|R| + \|R\| \times |S|$

Optimized Approach

- Start merging as soon as sorted runs from R and S can fit into memory
 - $B > N(R, i) + N(S, j)$, $i, j = \#$ passes from sorting R, S
- Create sorted runs of R and merge partially to get (R, i)
- Create sorted runs of S and merge partially to get (S, j)
- Merge (R, i) and (S, j)
- Analysis:** If $B > \sqrt{2|S|}$
 - # initial sorted runs of $S < \sqrt{\frac{|S|}{2}}$
 - Total # initial sorted runs of R and $S < \sqrt{2|S|}$
 - 1 pass is sufficient to merge and join the initial sorted runs
 - Total cost = $2(|R| + |S|) + (|R| + |S|) = 3(|R| + |S|)$

Grace Hash Join

- Assume R = build relation, S = probe relation
- Partition R into R_1, \dots, R_{B-1} by h
- Partition S using same h into S_1, \dots, S_{B-1}
- For each R_i , **build hashtable**
 - Allocate 1 buffer page for input, 1 page for output and $B - 2$ pages for hashtable T
 - For each tuple r in R_i , read it into input buffer and hash it into T using h' , $h \neq h'$
 - For each tuple s in S_i , read it into input buffer and **probe** T : If s matches with any r in bucket $h'(s)$, write $r \bowtie s$ into output buffer

Analysis

- By UHA: $|R_i| = \frac{|R|}{B-1}$. Let size of T be $f \times \frac{|R|}{B-1}$, f = fudge factor
 - Hence, $B > \frac{f|R|}{B-1} + 2 \approx \sqrt{f|R|}$ (1 input buffer, 1 output buffer) to prevent partition overflow
- Assuming no partition overflow:
 - Partitioning = $2(|R| + |S|)$
 - Probing = $|R| + |S|$
 - Read each page of R_i to build T
 - Read each page of S_i to probe
 - Total = $3(|R| + |S|)$

Query Evaluation: Misc Operations

Set Operations

- Intersection** $R \cap S$
 - Join with join predicate involving all columns of R and S
- Cross Product** $R \times S$
 - Join with join predicate = true (trivial)

Union $R \cup S$

- Sorting approach:
 - Sort R , sort S (on all attrs)
 - Combine R and S and removing duplicates
- Hashing approach: \approx Grace Hash Join
 - Partition (on all attrs) R into R_1, \dots, R_{B-1} ,
 - Partition S into S_1, \dots, S_{B-1}
 - Build hash table T_i for each R_i (suppose R is build relation)
 - For each $t \in S_i$, probe T_i and insert only if t not in T_i
- Difference** $R - S$
 - Sorting approach: \approx Sort-Merge Join using R as outer relation
 - Sort R , sort S (on all attrs)
 - Remove $t \in R$ if $t \in S$
 - Hashing approach: \approx Grace Hash, using R as build relation
 - Partition (on all attrs) R into R_1, \dots, R_{B-1} ,
 - Partition S into S_1, \dots, S_{B-1}
 - Build hash table T_i for each R_i
 - For each $t \in S_i$, probe T_i and discard t from T_i if t in T_i

Aggregate Operations

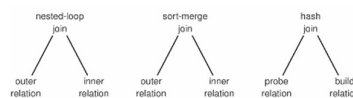
- Simple Aggregation:** Maintain running info while scanning table
 - Valid for SUM, COUNT, AVG, MIN, MAX
- Group-by Aggregation**
 - Sorting approach
 - Sort relation by 'GROUP BY' attributes
 - Scan relation and compute aggregate for each group
 - Hashing approach
 - Scan relation to build hash table on 'GROUP BY' attributes
 - Maintain running info for each group
- Using Index**
 - Use index I over table scan if I is a covering index for aggregation operation

Query Evaluation Approaches

- Materialized evaluation**
 - An operator is evaluated only when all of its operands has been completely evaluated/materialized
 - Materialize intermediate results to disk
- Pipelined evaluation**
 - Pass the output directly to its parent operator (no materialize)
 - Execution of operators is interleaved
 - Blocking operator: Operator that is unable to produce any output until it has received all the tuples from its child operators
- Iterator Interface of Pipelined evaluation**
 - open – initialization; allocates resources and operators' args
 - getNext – generates next output tuple/null if no more output
 - close: deallocate state information

Query Optimization

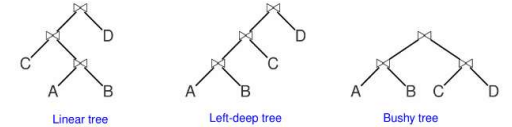
- Join Plan Notation:



Commutative 1.1. $R \times S \equiv S \times R$ 1.2. $R \bowtie S \equiv S \bowtie R$	Commutating σ with π 4.1. $\pi_L(\sigma_p(R)) \equiv \pi_L(\sigma_p(\pi_{L \setminus \text{attr}(p)}(R)))$
Associative 2.1. $(R \times S) \times T \equiv R \times (S \times T)$ 2.2. $(R \bowtie S) \bowtie T \equiv R \bowtie (S \bowtie T)$	Commutating σ with \times/\bowtie 5.1. $\sigma_p(R \times S) \equiv \sigma_p(R) \times S$, $\text{attr}(p) \subseteq \text{attr}(R)$ 5.2. $\sigma_p(R \bowtie_q S) \equiv \sigma_p(R) \bowtie_q S$, $\text{attr}(p) \subseteq \text{attr}(R)$ 5.3. $\sigma_p(R \cup S) \equiv \sigma_p(R) \cup \sigma_p(S)$
Idempotence 3.1. $\pi_A(\pi_B(R)) \equiv \pi_A(R)$, $A \subseteq B \subseteq \text{attr}(R)$ 3.2. $\sigma_{p_1}(\sigma_{p_2}(R)) \equiv \sigma_{p_1 \wedge p_2}(R)$	Commutating π with \times/\bowtie 6.1. $\pi_L(R \times S) \equiv \pi_{L_R}(R) \times \pi_{L_S}(S)$ 6.2. $\pi_L(R \bowtie_q S) \equiv \pi_{L_R}(R) \bowtie_q \pi_{L_S}(S)$, $\text{attr}(p) \cap \text{attr}(R) \subseteq L_R \ \& \ \text{attr}(p) \cap \text{attr}(S) \subseteq L_S$ 6.3. $\pi_L(R \cup S) \equiv \pi_{L_R}(R) \cup \pi_{L_S}(S)$

Query Plan Trees:

- Linear: At least 1 operand for each join operation is a base relation
- Bushy: There is a join operation where no operand is a base relation
- Left-deep: Every right join operand is a base relation
- Right-deep: Every left join operand is a base relation



Query Plan Enumeration

- Uses a bottom-up Dynamic Programming approach starting with size-1 relations and memoizing the best plan
- For every possible combination S of R_i , for every possible pair of partitions S_i and S_j in S , find the $\text{optPlan}(S)$ of joining S_i and S_j using the memoized $\text{optPlan}(S_i)$ and $\text{optPlan}(S_j)$

Input: A SQL query q on relations R_1, R_2, \dots, R_n
 Output: An optimal query plan for q

```

01. for  $i = 1$  to  $n$  do
02.    $\text{optPlan}(\{R_i\}) = \text{best access plan for } R_i$ 
03. for  $i = 2$  to  $n$  do
04.   for each  $S \subseteq \{R_1, \dots, R_n\}, |S| = i$  do
05.      $\text{bestPlan} = \text{dummy plan with cost}(\text{bestPlan}) = \infty$ 
06.     for each  $S_i, S_j \in S, |S_i| \in \{1, i\}, S = S_i \cup S_j$  do
07.        $p = \text{best way to join } \text{optPlan}(S_i) \text{ and } \text{optPlan}(S_j)$ 
08.       if  $\text{cost}(p) < \text{cost}(\text{bestPlan})$  then
09.          $\text{bestPlan} = p$ 
10.    $\text{optPlan}(S) = \text{bestPlan}$ 
11. return  $\text{optPlan}(\{R_1, \dots, R_n\})$ 
  
```

System R Optimizer

- Uses an enhanced Dynamic Programming approach that also considers sort order of query plan's output
 - Maintains $\text{optPlan}(S_i, o_i)$, o_i = sort order of output by query plan wrt S_i
 - $o_i = \text{null}$ if output is unordered or a sequence of attributes
- Prunes search space:
 - Considers only left-deep query plans
 - Avoids cross-product query plans
 - Considers early selections and projections

Query Plan Cost Estimation

- Reduction factor $rf(t_i)$: fraction of tuples in e that satisfy t_i i.e. $rf(t_i) = \frac{\|\sigma_{t_i}(e)\|}{\|e\|}$

Assumptions		
Uniformity	Independence	Inclusion
Uniform distribution of attribute values	Independent distribution of values in diff attributes	For $R \bowtie_{A=B} S$, if $\ \pi_A(R)\ \leq \ \pi_B(S)\ $, then $\pi_A(R) \subseteq \pi_B(S)$
$rf(A = c) \approx \frac{1}{\ \pi_A(R)\ }$	$rf(t_i \wedge t_j) \approx rf(t_i) \times rf(t_j)$	$rf(R.A = S.B) \approx \frac{1}{\max\{\ \pi_A(R)\ , \ \pi_B(S)\ \}}$

Equiwidth Histogram

- Each bucket has almost equal number of values
- All buckets have the same width/range size of B
- $\|\sigma_{A=c}(R)\| = \frac{\|b_i\|}{B}$
- $\|\sigma_{A \in [x,y]}(R)\| = \frac{f_1 \|b_i\|}{B} + \|b_{i+1}\| + \dots + \frac{f_2 \|b_{i+k}\|}{B}$

Equidepth Histogram

- Each bucket has almost equal number of tuples. Let B_i denote the width of bucket b_i
- A value can be contained in multiple buckets
- $\|\sigma_{A=c}(R)\| = \frac{f_1 \|b_i\|}{B_i} + \frac{f_2 \|b_{i+1}\|}{B_{i+1}} + \dots + \frac{f_k \|b_{i+k}\|}{B_{i+k}}$, for all b_j containing c
- $\|\sigma_{A \in [x,y]}(R)\| = \frac{f_1 \|b_i\|}{B_i} + \frac{f_2 \|b_{i+1}\|}{B_{i+1}} + \dots + \frac{f_k \|b_{i+k}\|}{B_{i+k}}$

Histogram with MCV

- Separately keep track of the frequencies of the top-k most common values and exclude them from the histogram

