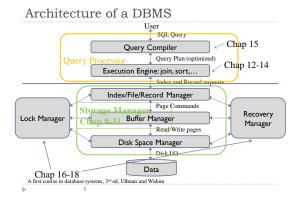


Slides based on "Database Management Systems" 3rd ed, Ramakrishnan and Gehrke

Coverage

- Text, chapters 8 through 15 (hw1 hw4)
- PKs, FKs, E-R to Relational: Text, Sec. 3.2-3.5, to pg. 77 inclusive, createdb.sql
- Basics of RAID: Sec. 9.2, Slides of Lecture 3
- SQL for creating and dropping tables (standardized), Not standardized: create indexes, commands for bulk loading big tables (Oracle and mysql cases).



Disks

Accessing a Disk Block

> Time to access (read/write) a disk block:

- seek time (moving arms to position disk head on track)
- rotational delay (waiting for block to rotate under head)
- transfer time (actually moving data to/from disk surface)

Seek time and rotational delay dominate for up to about IMB transfers, and DB pages are smaller than that

- Seek time varies from about 1 to 20msec
- Rotational delay varies from 0 to 10msec
- > Transfer rate is about Imsec per 4KB page

Key to lower I/O cost: reduce seek/rotation delays!

blocks on same track, followed by
 blocks on same cylinder, followed by
 blocks on adjacent cylinder

Arranging Pages on Disk

Next' block concept:

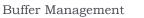
- > Logical block numbers of current disks follow this sequence
- Blocks that are accessed together frequently should be sequential on disk (by `next'), to minimize access time
- Use newly-initialized file systems for DB files to avoid OS file fragmentation
- For a sequential scan, <u>pre-fetching</u> several pages at a time is a big win!

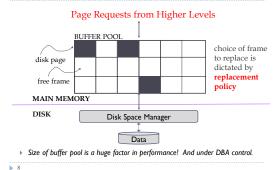
▶ 6

Important RAID Levels

Level 0: Striping but no redundancy

- Maximum transfer rate = aggregate bandwidth
- Stripe size can be many blocks, example 256KB
- With N data disks, read/write bandwidth improves up to N times
- ▶ Level I: Mirroring ← strongly recommended for redo log files
- Each data disk has a mirror image (check disk) Parallel reads possible, but a write involves both disks
- Level 0+1: Striping and Mirroring (AKA RAID 10) Maximum transfer rate = aggregate bandwidth
- With N data disks, read bandwidth improves up to N times
- Level 5: Block-Interleaved Distributed Parity (in wide use)
- Every disk acts as data disk for some blocks, and check disk for other blocks
- Most popular of the higher RAID levels (over 0+1). Dbs3 has RAID 5, even for redo log file (so not best performance for actions that change the database)





File Organization

▶ 7

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1. Unsorted, or heap file

Records stored in random order

2. Sorted according to set of attributes

- ▶ E.g., file sorted on <age>
- Or on the combination of <age, salary>

No single organization is best for all operations

- E.g., sorted file is good for range queries
- But it is expensive to insert records
- We need to understand trade-offs of various organizations

Heap

Unordered Files: Heap

- simplest file structure
- contains records in no particular order
- as file grows and shrinks, disk pages are allocated and deallocated

> To support record level operations, we must:

- keep track of the pages in a file
- keep track of free space on pages
- keep track of the records on a page

10

Data Organization

- Index/File/Record Manger provides abstraction of file of records (or short, file)
- File of records is collection of pages containing records
- A File can be a heap table, a heap table accessed via a certain index, a sorted table, or a certain index
- File operations
 - read/delete/modify a record (specified using record id)
 - insert record
 - scan all records, search with equality selection, search with range selection

Record id functions as data locator

- > contains information on the address of the record on disk
- e.g., page and record offset in page
- "search-by-address"

▶ 11

QP to Storage Engine API

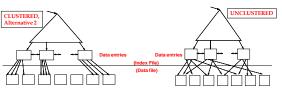
- Storage Engine works on one "File" at a time, that is, in one call from the QP, which could be in the middle of doing a join of two tables, or a sort, or ...
- > Table scan and index scan are just scans of two kinds of Files
- Cost models are based on the costs of the various calls into the Storage Engine, since it does all the disk i/o.
- See Figure 8.4 for various costs.

Indexing, starts in Chap. 8, then continues in 10 and 11

Alternatives for Data Entry k^* in Index (pg. 276 in Chap 8)

Data record with key value k
 Leaf node stores actual record
 Only one such index can be used (without replication)
 <k, rid> rid of data record with search key value k
 Only a pointer (rid) to the page and record are stored
 <k, list of rids> list of rids of records with search key k
 Similar to previous method, but more compact
 Disadvantage is that data entry is of variable length
 Don't worry about this case for exams
 Several indexes with alternatives 2 and 3 may exist

Clustered vs. Unclustered Index



Clustered index: order of data records is close to the sort order

Dynamic data structure (as opposed to ISAM)

Height is log F N (F = fanout, N = # leaf pages)

Each node (except root) contains d <= m <= 2d entries</p>

> But insert/delete more complex due to occupancy constraint

> Insert/delete may trigger re-structuring at all levels of tree

Minimum 50% occupancy constraint

Parameter d is called the order of the tree

- Here: loaded from ordered data, so records fall in order naturally.
- However, the most common kind of clustered index uses Alternative 1, not Alternative 2 as shown above, see next slide for picture
- Unclustered: must be Alternative 2 (or 3, but we're not worrying about that case)

▶ 15

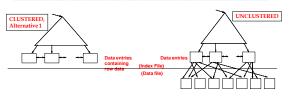
B+ Tree

Most Widely Used Index

Tree is height-balanced

Search just like in ISAM

Clustered vs. Unclustered Indexes



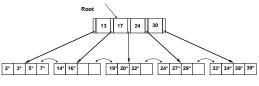
- > Clustered index: order of data records is close to the sort order
- The most common kind of clustered index uses Alternative 1, as shown above
- If see "clustered index" without Alternative specified, assume Alternative 1.

16

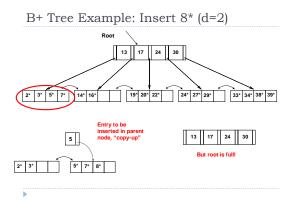
▶ 14

B+ Tree Example

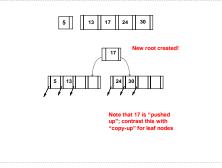
> Search begins at root, key comparisons direct it to a leaf



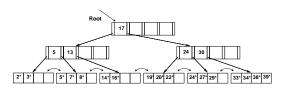
Based on the search for 15*, we know it is not in the tree!



B+ Tree Example: Insert 8* (d=2)



Example B+ Tree After Inserting 8*



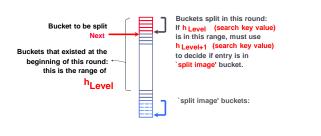
Root was split, leading to increase in height

We can avoid split by re-distributing entries, but his is usually not done in practice for insertions

Linear Hashing

- > Dynamic hashing scheme
- > Handles the problem of long overflow chains
 - But does not require a directory!
 - Deals well with collisions!
- Main Idea: use a family of hash functions **h**₀, **h**₁, **h**₂, ...
 - h_i(key) = h(key) mod(2ⁱN)
 N = initial number of buckets
 - If N = 2^{d_0} , for some d_0 , \mathbf{h}_i consists of applying \mathbf{h} and looking at the last d_i bits, where $d_i = d_0 + i$
 - \mathbf{h}_{i+1} doubles the range of \mathbf{h}_i (similar to directory doubling)

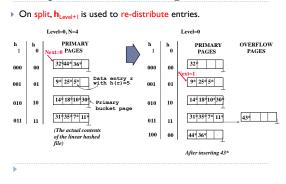
Overview of Linear Hashing



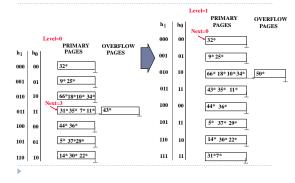
Linear Hashing Properties

- Directory avoided in LH by using overflow pages
- Buckets are split round-robin
 - Splitting proceeds in `rounds'
 - Round ends when all N_R initial buckets are split (for round R)
 - Buckets 0 to Next-1 have been split; Next to N_R yet to be split.
 - Current round number referred to as Level
- Search for data entry r :
 - If $\mathbf{h}_{Level}(r)$ in range `Next to N_R ', search bucket $\mathbf{h}_{Level}(r)$
 - Otherwise, apply $\mathbf{h}_{Level+1}(r)$ to find bucket

Example of Linear Hashing



End of a Round



Cost of Operations

	(a) Scan	(b) Equality	(c) Range	(d) Insert	(e) Delete
(1) Heap	BD	0.5BD	BD	2D	Search +D
(2) Sorted	BD	Dlog 2B	D(log 2 B + # pgs with match recs)	Search + BD	Search +BD
(3) Clustered	1.5BD	Dlog f 1.5B	D(log F 1.5B + # pgs w. match recs)	Search + D	Search +D
(4) Unclust. Tree index	BD(R+0.15)	D(1 + log f 0.15B)	· · ·	Search + 2D	Search + 2D
(5) Unclust. Hash index	BD(R+0.125)	2D	BD	Search + 2D	Search + 2D

> 27

Notes on these costs

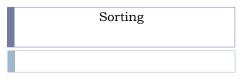
- B = # pages of data in file
- D = time for one (random) i/o
- R = # records/page
- Without further info on index size, use 10% table size
- Here, scan of clustered table takes 1.5 BD, where B = #pages data would take in a heap file, but if B = no of data pages of Btree, average of 2/3 full, then should be just BD.
- Similarly, scan of FILE by unclustered tree index = .1 (size of table) + RBD
- Later, we said Size(Clustered index) = I.I (Size of data), which really only should be used if the B-tree nodes are full.
- > B-tree nodes can be full after bulk load from sorted data.

General External Merge Sort

- ▶ To sort a file with N pages using B buffer pages:
 - ▶ Pass 0: use *B* buffer pages. Produce $\lceil N / B \rceil$ sorted runs of *B* pages each.
 - Pass 2, ..., etc.: merge B-1 runs.

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	INPUT 1 INPUT 2 OUTPUT	
Disk	B Main memory buffers	Disk



Cost of External Merge Sort

- Number of passes: $1 + \lceil \log_{B-1} \lceil N / B \rceil \rceil$
- Cost = 2N * (# of passes), assuming we need to read the input from a FILE and write the output to a FILE.
- Can save some i/o using pipelining in and out.
- Example: with 5 buffer pages, sort 108 page file:
 - Pass 0: ceil(108/5) = 22 sorted runs of 5 pages each (last run is only 3 pages)
 - ▶ Pass I: [22 / 4]= 6 sorted runs of 20 pages each (last run is only 8 pages)
 - Pass 2: 2 sorted runs, 80 pages and 28 pages
 - Pass 3: Sorted file of 108 pages

Query Evaluation

Executing Selections

- Find the most selective access path, retrieve tuples using it > Then, apply any remaining terms that don't match the index
- Most selective access path: index or file scan estimated to require the fewest page I/Os
 - Consider day<8/9/94 AND bid=5 AND sid=3</p>
- > If we have B+ tree index on day, use that access path
 - Then, bid=5 and sid=3 must be checked for each retrieved tuple
 - day condition is primary conjunct
- Alternatively, use hash index on <bid, sid> first
 - Then, day<8/9/94 must then be checked</p>

Example of matching indexes

Pg. 399: fix error Sailors→ Reserves on line 8

Reserves (<u>sid: integer, bid: integer, day: dates</u>, rname: string) ← rname column added here with indexes:

- Index1: Hash index on (rname, bid, sid) Matches: rname='Joe' and bid = 5 and sid=3
- Doesn't match: rname='Joe' and bid = 5
- Index2:Tree index on (rname, bid, sid)
 - Matches: rname='Joe' and bid = 5 and sid=3
 - Matches: rname='joe' and bid = 5, also rname = 'joe'
 - Doesn't match: bid = 5
- Index3:Tree index on (rname)
- Index4: Hash index on (rname)
- These two match any conjunct with rname='Joe' in it

Using an Index for Selections

Cost influenced by:

- Number of qualifying tuples
- Whether the index is **clustered** or not
- Cost of finding qualifying data entries is typically small
- ▶ E.g., SELECT *
 - FROM Reserves R WHERE R.rname < 'C%'
- Assuming uniform distribution of names, 10% of tuples gualify, that is 10000 tuples
 - With a clustered index, cost is little more 100 I/Os
 - If not clustered, up to I 0K I/Os!

Hw4 Problem 2 (15.2, question 1) NPages(R) = 10000, 8000 usable bytes/page, 20 bytes/data entry

NTuples(R) = 800,000, so 800,000 *20 bytes/secondary index, = 2000 pgs Reduction Factor (RF) = 0.1

NPages(UI) = 2000 For unclustered indexes,

Cost(UI) = (NPages(UI)+NTuples(R)) * product of RF's of matching predicates For clustered indexes, the rows lie in the leaf pages, 2000 of them, and the level above that has 2000 data entries, or 20*2000 bytes = 25 pages. The level above that is the

- Alt. I: NPages(CI) = NLeafPages + NIndexPages = 10,000 + 25 = 10,025. Alt. 2: NPages(Cl) = index size + table size = 2000 + 10,000 = 12,000
- Cost(CI) = NPages(CI) * product of RF's of matching predicates

Index #I: Unclustered hash index on eid

Index #2: Unclustered B+ Tree index on sal

Index #3: Unclustered hash index on age

- Index #4: Clustered B+ Tree index on <age, sal> See the hw4 solution for further info on this.

Projection with Sorting

- Modify Pass 0 of external sort to eliminate unwanted fields
 - Runs of about 2B pages are produced
- Tuples in runs are smaller than input tuples
- > Size ratio depends on number and size of fields that are dropped
- Modify merging passes to eliminate duplicates
 - > Thus, number of result tuples smaller than input
- Difference depends on number of duplicates
- Cost
 - In Pass 0, read original relation (size M), write out same number of smaller tuples
 - In merging passes, fewer tuples written out in each pass. Using Reserves example, 1000 input pages reduced to 250 in Pass 0 if size ratio is 0.25

Projection with Hashing

Partitioning phase:

- Read R using one input buffer. For each tuple, discard unwanted
- fields, apply hash function *h1* to choose one of **B-1** output buffers > Result is **B-1** partitions (of tuples with no unwanted fields), tuples
- from different partitions guaranteed to be distinct

Duplicate elimination phase:

- For each partition, read it and build an in-memory hash table, using hash h2 on all fields, while discarding duplicates
- If partition does not fit in memory, can apply hash-based projection algorithm recursively to this partition

Cost

 Read R, write out each tuple, but fewer fields. Result read in next phase

Discussion of Projection

- Sort-based approach is the standard
- better handling of skew and result is sorted.If index on relation contains all wanted attributes in its
- search key, do *index-only* scan
 - Apply projection techniques to data entries (much smaller!)
- If an ordered (i.e., tree) index contains all wanted attributes as prefix of search key, can do even better:
 - Retrieve data entries in order (index-only scan)
 - Discard unwanted fields, compare adjacent tuples to check for duplicates

Simple Nested Loops Join

foreach tuple r in R do foreach tuple s in S do if $r_i == s_i$ then add <r, s> to result

- For each tuple in the *outer* relation R, we scan the entire *inner* relation S.
- Cost: M + p_R * M * N = 1000 + 100*1000*500 I/Os

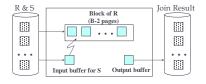
Page-oriented Nested Loops join:

- > For each page of R, get each page of S, and write out matching pairs
- Cost: M + M*N = 1000 + 1000*500
- If smaller relation (S) is outer, cost = 500 + 500*1000

Block Nested Loops Join

one page input buffer for scanning the inner S

- one page as the output buffer
- remaining pages to hold ``block" of outer R
- For each matching tuple r in R-block, s in S-page, add <r, s> to result. Then read next R-block, scan S, etc.



Example of NLJ

SELECT * FROM Reserves R1, Sailors S1 WHERE R1.sid=S1.sid

Reserves:

- 40 bytes long tuple, 100K records, 100 tuples per page, 1000 pages, suppose B+ Tree index on bid, another on sid
- Sailors:
 - 50 bytes long tuple, 40K tuples, 80 tuples per page, 500 pages, suppose B+ tree index on sid

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Examples of Block Nested Loops

Cost: Scan of outer + #outer blocks * scan of inner #outer blocks =

[# of pages of outer / blocksize]

- With Reserves (R) as outer, and 100 pages of R:
 - Cost of scanning R is 1000 I/Os; a total of 10 blocks.
 - Per block of R, we scan Sailors (S); 10*500 I/Os. Total 6000 i/os
- With 100-page block of Sailors as outer:
 - Cost of scanning S is 500 I/Os; a total of 5 blocks. Per block of S, we scan Reserves; 5*1000 I/Os.
 - Total 5500 I/Os
- With 50-page block of S as outer (hw4 #1 part 2, S has 200 pages, R has 1000)
- Cost of scanning S is 200 I/Os; a total of 4 blocks.
- Per block of S, we scan Reserves; 4*1000 I/Os. Total 4,200 I/Os

Executing Joins: Index Nested Loops

foreach tuple r in R do for each tuple s in S where $r_i == s_j do$ add <r, s> to result Cost = M + (M*p_R) * (cost of finding matching S tuples) M = number of pages of R, p_R = number of R tuples per page

- If relation has index on join attribute, make it inner relation
- For each outer tuple, cost of probing inner index is 1.2 for hash index, 2-4 (say 2 for simplicity) for B+, plus cost to retrieve matching S tuples Clustered index:
 - Alt I: typically no more I/Os (data entry has whole row)
- Alt 2: typically single I/O (data entry has RIDs, but target rows are clustered in table)
- Unclustered index I I/O per matching S tuple

Example of Index Nested Loops (1/2)

Case I: B+tree-index on sid of Sailors

- Choose Sailors as inner relation
- Scan Reserves: 100K tuples, 1000 page I/Os
- For each Reserves tuple
 - > 2 I/Os to get data entry in index (simplifying 2-4 in text)
 - No more i/o if clustered Alt 1, 1 I/O to get (the exactly one) matching Sailors tuple (primary key), clustered Alt 2 or unclustered
- Total: 201,000 or 301,000 I/Os, terrible. 3010 s = 50 min.

Example of Index Nested Loops (2/2)

Case 2: B+ tree-index on sid of Reserves

- Choose Reserves as inner
- Scan Sailors: 40K tuples, 500 page I/Os

For each Sailors tuple

- > 2 I/Os to find index page with data entries (simplified from 2-4)
- > Assuming uniform distribution, 2.5 matching records per sailor
- Cost of retrieving records is nothing (Alt I clustered), single I/O (Alt. 2 clustered index) or 2.5 I/Os (unclustered index)
- Total: 80,500 I/Os (clustered Alt I), 120,500 I/Os (clustered Alt 2) or 180,500 I/Os (unclustered) All bad.
- Better to use block NLJ here, if required to do NLJ.

Sort-Merge Join

- Sort R and S on the join column
- > Then scan them to do a merge on join column:
 - > Advance scan of R until current R-tuple >= current S tuple
 - > Then, advance scan of S until current S-tuple >= current R tuple
 - Repeat until current R tuple = current S tuple
 - > At this point, all R tuples with same value in Ri (current R group) and all S tuples with same value in Sj (current S group) match
 - Output <r, s> for all pairs of such tuples
 - Resume scanning R and S

Sort-Merge Join Cost

R is scanned once

1

- Each S group is scanned once per matching R tuple
- > Multiple scans per group needed only if S records with same join attribute value span multiple pages
- Multiple scans of an S group are likely to find needed pages in buffer
- Cost: (assume B buffers)
 - 2M (1+log B-1 (M/B)) + 2N (1+log B-1 (N/B)) + (M+N)
- The cost of scanning, M+N, could be M*N worst case (very unlikely!)
- In many cases, join attribute is primary key in one of the tables!

Sort-Merge Join Cost: hw4 #1, part 3

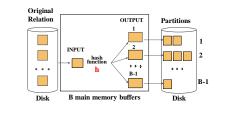
- R JOIN S on R.a = S.b, where b is the PK of S. B=1500.
- I. Sort R and S on the join column:
- Sort R on a: 100K pages.
 - S on b: 20K pages Pass 0: 100K/1500 = 68 runs of 1500 pages; 20K/1500 = 14 Pass 1: merge 68 runs into one merge 14 into one
- So both are 2-pass sorts, unlike original-problem setup
- Cost = 2*2*M+2*2*N = 4*(100K+20K) = 480K i/os.
- > 2. Then scan them to do a merge on join column:
- R is scanned once, each row matching one row of S Cost = M+N reads (ignore output costs by problem) = 120K i/os
- Total cost = 600K i/os (not 100x old answer!)
- Note: this is not using the optimization which yields 3(M+N) for 2-pass sorts: by that algorithm, cost = 3*120K = 360K i/os.

2-Pass Sort-Merge Join

- With enough buffers, sort can be done in 2 passes
- First pass generates N/B sorted runs of B pages each If one page from each run + output buffer fits in memory, then merge can be done in one pass; denote larger relation by L
- $2L/B + I \le B$, holds if (approx) $B > \sqrt{2L}$
- One optimization of sort allows runs of 2B on average
- First pass generates N/2B sorted runs of 2B pages each
- Condition above for 2-pass sort becomes B $> \sqrt{L}$
- Merge can be combined with filtering of matching tuples The cost of sort-merge join becomes 3(M+N)

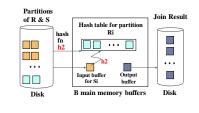
Hash-Join: Partitioning Phase

- Partition both relations using hash function h
- R tuples in partition i will only match S tuples in partition i



Hash-Join: Probing Phase

Read in a partition of R, hash it using h2 (<> h!) Scan matching partition of S, search for matches.



Hash-Join Properties

- #partitions $k \le B-I$ because one buffer is needed for scanning input
- Assuming uniformly sized partitions, and maximizing k:
- ▶ k= B-1, and M/(B-1) <= B-2, i.e., $B > \sqrt{M}$
- M is smaller of the two relations!
- If we build an in-memory hash table to speed up the matching of tuples, slightly more memory is needed
- If the hash function does not partition uniformly, one or more R partitions may not fit in memory
- Can apply hash-join technique recursively to do the join of this Rpartition with corresponding S-partition.

Cost of Hash-Join

- In partitioning phase, read+write both R and S: 2(M+N)
- In matching phase, read both R and S: M+N
- So Cost with partitioning = 3(M+N)
 - With sizes of 1000 and 500 pages, total is 4500 I/Os
 - (not counting materialization of result)
- If hash table of one table's data fits in memory, cost = M+N
- > If hash table for a partition doesn't fit in memory, cost exceeds above estimate.

Hash-Join vs Sort-Merge Join

- \triangleright Given sufficient amount of memory both have a cost of 3(M+N) I/Os
- Hash Join superior on this count if relation sizes differ greatly
- Hash Join shown to be highly parallelizable
- > Sort-Merge less sensitive to data skew, and result is sorted

Query Optimization: Chap. 15

CS634 Lecture 12, Mar 9, 2016

Slides based on "Database Management Systems" 3rd ed, Ramakrishnan and Gehrke

Block Optimization

- Block = Unit of optimization
- For each block, consider:
 - 1. All available access methods, for each relation in FROM clause
 - 2. All left-deep join trees
 - all ways to join the relations one-at-a-time
 - all relation permutations and join methods
- Recall:
- Left table = outer table of a nested loop join
- Left table of NLJ can be pipelined: rows used one at a time in order
- But need to consider other join methods too, giving up pipelining in many cases

$\sigma\pi \times \text{Expressions}$

- Query is simplified to a selection-projection-cross product expression
 - Aggregation and grouping can be done afterwards
- Optimization with respect to such expressions
- Cross-product includes conceptually joins
 Will talk about equivalences in a bit

Size Estimation and Reduction Factors

SELECT attribute list FROM relation list WHERE term₁ AND ... AND term_k

- Maximum number of tuples is cardinality of cross product
- Reduction factor (RF) associated with each term reflects its
- impact in reducing result size
- Implicit assumption that terms are independent!
- > col = value has RF =1/NKeys(l), given index l on col
- col1 = col2 has RF = 1/max(NKeys(l1), NKeys(l2))
- > col > value has RF = (High(I)-value)/(High(I)-Low(I))

- Single-Relation Plans
- FROM clause contains single relation
- Query is combination of selection, projection, and aggregates (possibly GROUP BY and HAVING, but these come late in the logical progression, so usually less crucial to planning)
- Main issue is to select best from all available access paths (either file scan or index)
- Access path involves the table and the WHERE clause
- Another factor is whether the output must be sorted
 E.g., GROUP BY requires sorting
 - Sorting may be done as separate step, or using an index if an indexed access path is available

Plans Without Indexes

- Only access path is file scan
- Apply selection and projection to each retrieved tuple
- Projection may or may not use duplicate elimination, depending on whether there is a DISTINCT keyword present
- GROUP BY:
- Write out intermediate relation after selection/projection
- (or pipeline into sort)
- Sort intermediate relation to create groups
- Apply aggregates on-the-fly per each group
- > HAVING also performed on-the-fly, no additional I/O needed

Plans With Indexes

There are four cases:

- 1. Single-index access path
- Each index offers an alternative access path
- Choose one with lowest I/O cost
- Non-primary conjuncts, projection, aggregates/grouping applied next
- 2. Multiple-index access path
 - Each index used to retrieve set of rids
 - Rid sets intersected, result sorted by page id
 - Retrieve each page only once
- Non-primary conjuncts, projection, aggregates/grouping applied next

Plans With Indexes (contd.)

3. Index-only access path

- If all attributes in query included in index, then there is no need to access data records: index-only scan
- > If index matches selection, even better: only part of index examined
- > Does not matter if index is clustered or not!
- > If GROUP BY attributes prefix of a tree index, no need to sort!
- > Example: With tree index on rating

SELECT max(rating),count(*) FROM Sailors S

Note count(*) doesn't require access to row, just RID.

Plans With Indexes (contd.)

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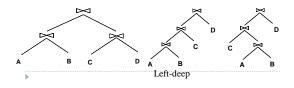
SELECT max(rating),count(*) FROM Sailors S

Note count(*) doesn't require access to row, just RID.

Queries Over Multiple Relations

In System R only left-deep join trees are considered

- In order to restrict the search space
- Left-deep trees allow us to generate all fully pipelined plans
 - > Intermediate results not written to temporary files.
 - > Not all left-deep trees are fully pipelined (e.g., sort-merge join)



Enumeration of Left-Deep Plans

- > Among all left-deep plans, we need to determine:
 - the order of joining relations
 - the access method for each relation
 - the join method for each join
- Enumeration done in N passes (if N relations are joined):
 - Pass I: Find best 1-relation plan for each relation
 - Pass 2: Find best way to join result of each 1-relation plan (as outer) to another relation - result is the set of all 2-relation plans
- Pass N: Find best way to join result of a (N-1)-relation plan (as outer) to the N'th relation - result is the set of all N-relation plans
- Speed-up computation using dynamic programming