Advanced Databases

Stratis D. Viglas

University of Edinburgh

Course logistics

- Lecturer: Stratis Viglas
 - email: sviglas@inf.ed.ac.uk
- Days/Times: Mon & Thu, 11:10-12:00
- Office hours: Mon, Thu 12:00-13:00 (or, by appointment)
 - ▶ *Room*: IF. 5.11
- Course webpage: www.inf.ed.ac.uk/teaching/courses/adbs
- Mailing list: adbs-students@inf.ed.ac.uk

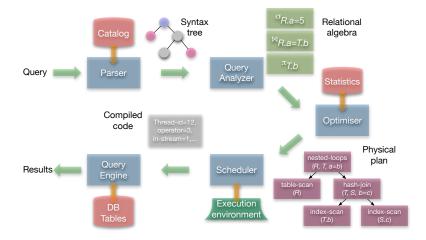
Syllabus

- Introduction
- Relational databases overview
 - Data model. evaluation model
- Storage
 - Indexes, multidimensional data
- Query *evaluation*
 - Join evaluation algorithms, execution models
- Query optimisation
 - Cost models, search space exploration, randomised optimisation
- Concurrency control and recovery
 - Locking and transaction processing
- Parallel databases

Assignments and software

- Programming assignments
- The attica database system
 - Home-grown *RDBMS*, written in Java
 - Visit inf.ed.ac.uk/teaching/courses/adbs/attica to download the system and the API documentation
 - All programming assignments will be using the attica front-end and code-base
- Plagiarism policy: You cheat, you're caught, you fail
 - No discussion

Query cycle



Three basic building blocks

- Attribute
 - ► A (name, value) *pair*
- Tuple
 - A set of attributes
- Relation
 - A set of tuples with the same schema

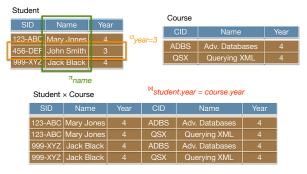


| SID | Name | Year |
|---------|------------|----------|
| 123-ABC | Mary Jones | 4 |

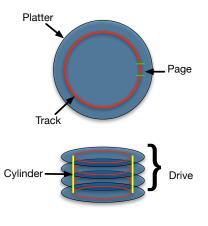
| SID | Name | Year |
|---------|------------|----------|
| 123-ABC | Mary Jones | 4 |
| 456-DEF | John Smith | 3 |
| | | |
| 999-XYZ | Jack Black | 4 |

Data manipulation

- Operations to *isolate* a *subset* of a *single relation*: Selection (σ), Projection (π)
- All set operations: Intersection, union, Cartesian product, set difference
- More *complex* operations: *Joins* (⋈), semi-joins, . . .



Data storage



- Disk drives are organised in records of 512 bytes
- The DB (and the OS) *I/O unit* is a *disk page* (typically, 4,096 bytes long)
- *Pages* (and records) are *stored* on *tracks*
- *Tracks* make up a *platter* (or a disk)
- *Platters* make up a *drive*
- The *same tracks* across all *platters* make up a *cylinder*
- The *disk head* (arm) reads the same block of all tracks on all platters

A bit of perspective

- The *dimensions* of the *head* are *impressive*¹. With a *width* of less than a *hundred nanometers* and a *thickness* of about *ten*, it flies above the platter at a *speed* of up to *15,000 RPM*, at a *height* that is the equivalent of *40 atoms*. If you start multiplying these infinitesimally small numbers, you begin to get an idea of their significance.
- Consider this little *comparison*: if the *read/write head* were a *Boeing* 747, and the *hard-disk platter* were the *surface of the Earth*
 - ► The *head* would *fly* at *Mach 800*
 - At less than one centimeter from the ground
 - And count every blade of grass
 - Making fewer than 10 unrecoverable counting errors in an area equivalent to all of *Ireland*

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¹Source: Matthieu Lamelot, Tom's Hardware.

What about flash memory and solid state?

• The *geometry* is different

There are no tracks, or platters, or cylinders or anything of the sort

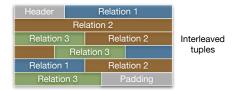
• But the *issues* are *similar*

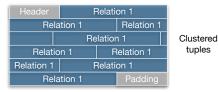
- Data is still accessed in *blocks*
- Blocks are still organised in pages
- Sequential vs. random I/O is still a problem
- Most of the things we say in this course are *applicable* to solid state as well
 - Added complexity: write/read asymmetry

Storing tuples

Every disk block contains

- ► A header
- Data (i.e., tuples)
- Padding (maybe)
- Two ways of storing tuples
 - Either interleave tuples of multiple relations, or
 - Keep the tuples of the same relation clustered





Advantages of clustering

- Scan a relation of X tuples, Y tuples per block
 - If unclustered, worst case scenario: read X blocks
 - Clustered: read X/Y blocks
- How about *clustering* disk *blocks*?
 - Reduces unnecessary arm movement

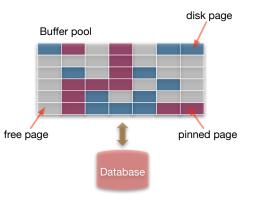


. .



The buffer manager

- Though the *data* is *on disk*, real *processing* is in *main memory*
- Disk blocks are read and put into the *buffer pool*
 - A collection of *memory* pages
- The *buffer manager* manages the buffer pool
 - Keeping track of page references, replacing pages if full, ...



What does the buffer manager do?

• When a *page is requested* it:

- Checks to see if the page is in the buffer pool; if so it returns it
- If not, it checks whether there is room in the buffer pool; if so it reads it in and places it in the available room
- If not, it picks a page for replacement; if the page has been "touched" it writes the page to disk and replaces it
- ► In all three cases, it *updates the reference count* for the requested page
- If necessary, it pins the new page
- It returns a handle to the new page

Page replacement

- Least recently used (LRU): check the number of references for each page; replace a page from the group with the lowest count (usually implemented with a priority queue)
 - Variant: clock replacement
- First In First Out (FIFO)
- Most recently used (MRU): the inverse of LRU
- Random!

Why not use the OS

- The OS implements virtual memory, so why not use it?
 - Page reference patterns and pre-fetching: the RDBMS in most cases knows which page will be accessed later (think of a clustered sequential scan)
 - Different page replacement policies according to the reference pattern (check p. 322 of your book)
 - ► Page pinning: certain pages should not be replaced
 - Control over when a page is written to disk: at times, pages need to be forced to disk (we'll revisit that when discussing crash recovery)

Indexing and sorting

- Can be summarised as:
 - Forget whatever you've learned about indexing, searching and sorting in main memory (well, almost ...)
- Remember, we are operating over disk files
 - The main idea is to minimise disk I/O and not number of comparisons (i.e., complexity)
 - ► Just an idea: comparing two values in memory costs 4.91 · 10⁻⁸ seconds; Comparing two values on disk costs 18.2 · 10⁻⁵ seconds (3 orders of magnitude more expensive.)

Indexing

Indexing functionality

- Indexes can be used for:
 - Lookup queries (e.g., [...] where value = ''foo'')
 - ▶ Range queries (e.g., [...] where value between 20 and 45)
 - Join processing (after all, predicates are value-based, aren't they?)
- The above uses, and much more, are what we call access methods

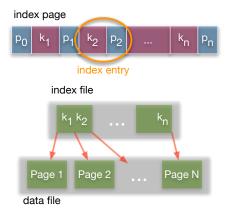
Two main classes

• Tree-structured indexes

- Much like you would use a binary tree to search, but with a higher key-per-node cardinality
- Retains order
- Great for *range queries*
- Both one-dimensional and multi-dimensional
- Hash-based indexes
 - Fully randomized (i.e., no order)
 - Great for single *lookup queries*

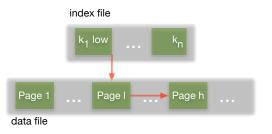
Sorted indexes

- The basic idea:
 - An *index* is on *an* (collection of) attribute(s) of a relation (called the *index key*)
 - It is much smaller than the relation
 - Index pages contain (key, pointer) pairs
 - ★ *key* of the *index*
 - ★ *pointer* to the *data page*
 - Plus one additional pointer (*low key*)



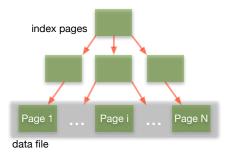
How does it answer range queries?

- Query is low ≤ value ≤ high
- Do a *binary search* on the *index file* to identify the *page containing the low key*
- *Keep scanning* the data file until the *high key* is *found*
- All done!

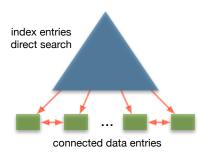


Potential problem (and the solution)

- The *index* is *much smaller* than the *relation*, but it's *still big*
- *Binary search* on it is *still expensive*
 - Remember, *data* is *on disk*
 - Have to access half the index file pages, plus the pages satisfying the predicate, all doing random I/O
- Why not build an *index on the index*?
 - ► Tree!

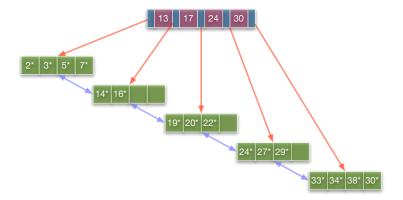


B+trees: the most widely used indexes



- Insertion/deletion at log_f N cost (f = fanout, N = # leaf pages)
- Tree is *height-balanced*
- Minimum 50% occupancy (except for root)
- Characterised by its order d; each node contains d ≤ m ≤ 2d entries
- *Equality* and *range* searches are *efficient*

B+tree example



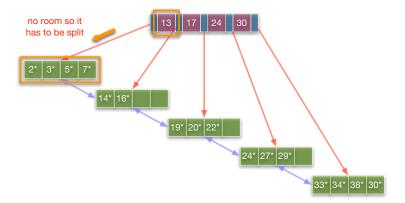
B+trees in practice

- Typical order: 100, typical fill-factor: 67%
 - Average fan-out: 133
- Typical capacities
 - Height 3: 2,532,637
 - Height 4: 312,900,700 (!)
- The top levels can often be kept in memory
 - Ist level: 4,096, or 8,192 bytes (1 page)
 - 2nd level: 0.5, or 1MB (133 pages)
 - 3rd level: 62, or 133MB

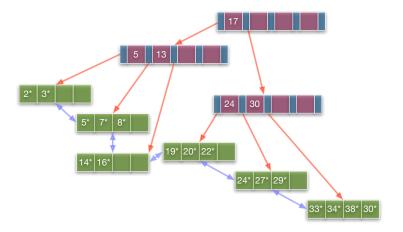
B+tree insertion

- Find correct leaf L
- Put data entry into L
 - If there is enough space in L, done!
 - If there is no space, L needs to be split into L and L'
 - Redistribute entries evenly in L and L'
 - Insert index entry pointing to L' into the parent of L
- Ascend the tree recursively, splitting and redistributing as needed
- *Tree tries to grow horizontally*; *worst case* scenario: a *root split* increases the height of the tree

B+tree insertion: 8*



B+tree insertion: 8*



Insertion observations

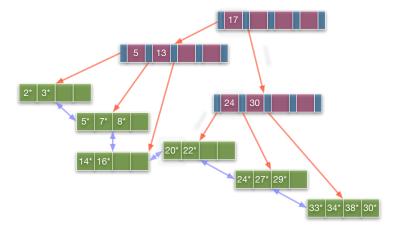
- Minimum occupancy is guaranteed at both leaf and non-leaf pages
- A *leaf split* leads to *copying* the key; a *non-leaf* split leads into *pushing up* the key (*why?*)
- The tree tries to *first grow horizontally* and if this is not possible, *then vertically*
 - In the example we could have avoided the extra level by redistributing
 - But in practice this is hardly ever done (why?)

B+tree deletion

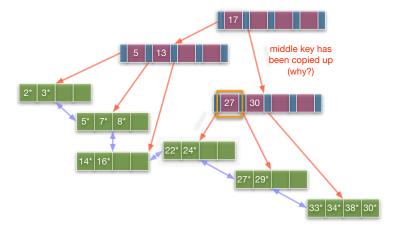
• Find leaf L where entry belongs

- *Remove* the entry
- If L is half-full, done!
- ▶ If *L* only has *d* − 1 entries
 - * Try to *redistribute* entries, *borrowing* from an *adjacent sibling* of L
 - * If redistribution fails, merge L and its sibling
 - ★ If merge has occurred, delete the entry for the merged page from the parent of L
- Ascend the tree recursively, performing the same algorithm
- Merge could propagate to the root, decreasing the trees height

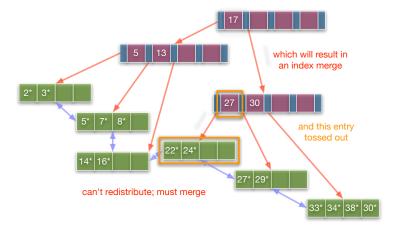
B+tree deletion: 19*



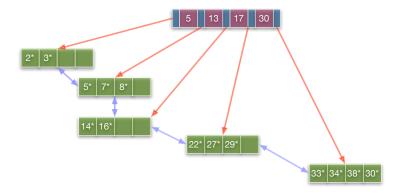
B+tree deletion: 20*



B+tree deletion: 24*



B+tree after deletion of 24^*



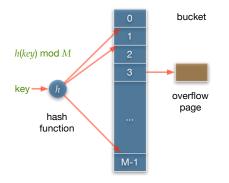
Summary of B+tree indexes

- Ideal for range searches, good for equality searches
- Highly *dynamic* structure
 - Insertions and deletions leave tree height-balanced, log_f N cost
 - For most typical implementations, height is rarely greater than 3 or 4, occupancy at 67%
 - Which means that the *index is almost always in memory*! (remember the buffer pool?)
 - Almost always better than maintaining a sorted file
 - The most optimised RDBMS structure

Hash indexes

- *Hash-based indexes* are *good* for *equality* selections, *not* for *range* selections
 - In fact, they cannot support range selections (why?)
- Static and dynamic techniques exist here as well
 - ► *Trade-offs* similar to those between ISAM and B+trees

Static hashing



- Number of primary pages fixed
 - Allocated sequentially, never de-allocated
 - Overflow pages if needed
- h(k) mod M = bucket to which data entry with key k belongs (M = number of buckets)

Static hashing observations

• The *buckets contain* the *actual data*!

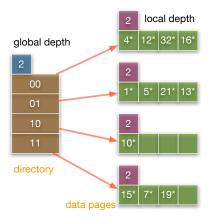
- But only the key is hashed
- No secondary index like in the tree case
- The *hash function must uniformly distribute* the *keys* across all buckets
 - Lots of ways to tune the hash function
- Again, *long overflow chains* of pages will develop, and pretty soon we're doing *random I/O*
 - ► *Need* a *dynamic* technique (big surprise here...)
 - Extendible hashing to the rescue

Extendible hashing

- Problem: bucket (i.e., primary page) becomes full
- Solution: re-organize the file by doubling the number of buckets
 - Are you crazy? Reading and writing out everything is expensive!
 - Why not keep a directory of buckets and double only the directory? Only read the bucket that overflowed
 - Directory much smaller; operation much cheaper

Extendible hashing example

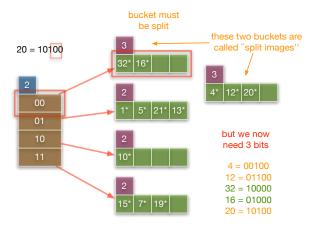
- Directory: array of size 4
- *Key k*, apply *hash function h(k)* and *translate* the result to *binary*
 - e.g., h(k) = 5 = 101
- Last global depth number of bits identify the bucket



Global, local depth and doubling

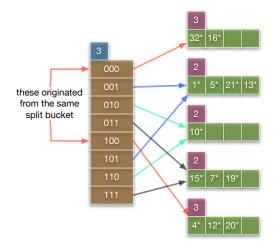
- *Global depth* (pertains to *directory*): maximum *number of bits* needed to tell which *bucket an entry belongs to*
- Local depth (pertains to bucket): maximum number of bits needed to tell whether an entry belongs to this bucket
- *Before* insertion (*local* = *global*) holds; *if insertion causes* (*local* > *global*) then *directory* needs to be *doubled*

Insertion example: h(k) = 20



so we must double the directory

Doubling the directory



Extendible hashing observations

- Directory fits in memory: equality search answered with only one disk I/O (two in the worst case!)
 - 100MB file, 100 bytes/tuple, 4kB pages, 1,000,000 data entries, 25,000 directory entries: fits in memory!
 - ► If the value distribution is skewed, directory grows large
 - Same hash-value entries are a problem (why?)
- Deletion: if removal empties bucket, then it can be merged with split image; if each directory entry points to the same bucket as its split image, the directory is halved

Linear hashing

- *Extendible hashing directory*: even if it is small, it is still a *materialised level of indirection*
- Though the *number of buckets grows linearly*, the size of the *directory* grows *exponentially*
- Objective: no directory, linear growth
- Linear hashing gets the job done

Why one, when you can have many?

- Key idea: *instead* of having a *single* hash function and using a *set of bits*, have *multiple hash functions*
 - Multiple hash functions implement the progressive doubling of the directory
- *Allocate* buckets *not* when they become *full*, *but* whenever we reach some *pretetermined load factor*
- Single bucket allocation
- Each bucket allocation results in another hash function to be used
- *Keep track* of the number of *buckets* and the number of *times* the number of buckets has *doubled*
- Discard unused hash functions

In more detail

- Use a *family* of *hash functions* h_0, h_1, h_2, \ldots
 - $h_i(key) = g(key) \mod (2^i M)$
 - *M* = *initial* number of *buckets*
 - g is some hash function (range is not [0, ..., N-1])
 - ▶ If $M = 2^{d_0}$, for some d_0 , h_i consists of applying g and looking at the last d_i bits, where $d_i = d_0 + i$.
 - h_{i+1} doubles the range of h_i (similar to directory doubling)

Bookkeeping

- Two variables: Next, and Level
 - N points to the *bucket* to be *split next*
 - L keeps track of the number of times the range of the hash function has doubled
- Splitting proceeds in 'rounds
 - ▶ Round ends when all M_R initial (for round R) buckets are split
 - Buckets 0 to N-1 have been split
 - Buckets N to M_R have yet to be split
- Current round is L

Search and insert

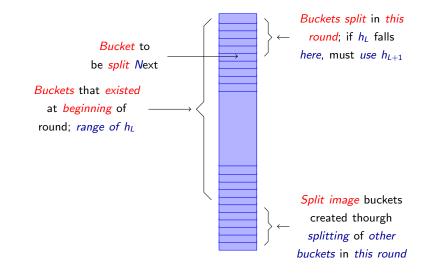
Search

- To find bucket for key K, find $h_L(K)$)
 - ▶ If $h_L(K) \in [N, ..., M_R]$, r belongs here
 - Else, *r* could belong to bucket $h_L(K)$ or bucket $h_L(r) + M_R$; we must apply $h_{L+1}(K)$ to find out.

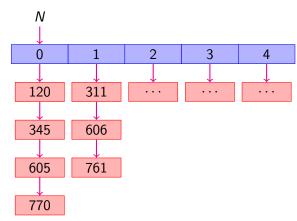
Insert

- Find bucket as above, by applying h_L or h_{L+1}
- If *bucket* to insert is *full*
 - Add overflow page and insert entry
 - (Maybe) Split bucket N and increment N

Linear hashing file



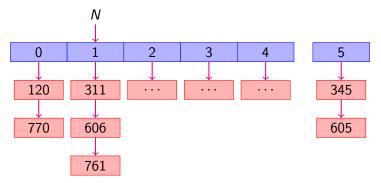
Splitting a bucket (0 in this case)



Hash functions

• $h_0(K) = K \mod 5$

Splitting a bucket (0 in this case)



Hash functions

•
$$h_1(K) = K \mod 10$$

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Algorithms in more detail

Lookup for key K *bucket* := $h_L(K)$; if *bucket* < N then *bucket* = $h_{L+1}(K)$

Expansion

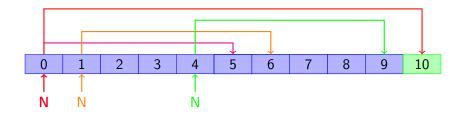
$$\begin{split} & \mathcal{N} := \mathcal{N} + 1; \\ & \text{if } \mathcal{N} = \mathcal{M} 2^L \text{ then} \\ & \mathcal{L} := \mathcal{L} + 1; \quad \mathcal{N} := 0; \end{split}$$

Contraction

N := N - 1;if N < 0 then L := L - 1; $N := M2^{L} - 1;$ The expansion process (round 0)

Expansion

N := N + 1;if $N = M2^{L}$ then $L := L + 1; \quad N := 0;$

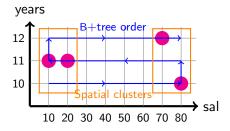


Linear hashing observations

- Can choose any criterion to trigger split
 - Typically, we want to maintain some load factor
- Since *buckets* are *split round-robin*, *long* overflow *chains do not develop!*
- Doubling of directory in extendible hashing is similar
 - Switching of hash functions is implicit in how the number of bits examined is increased

Why more than one dimensions?

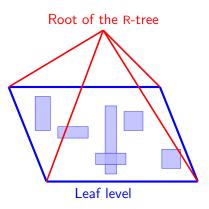
- Single-dimensional indexes are not enough
 - ► Consider a *composite search* key *e.g.*, an index on *(sal, years)*
 - The 2-dimensional space is linearised
 - We sort entries first by sal and then by years
- A multidimensional index clusters entries
 - Exploits nearness in multidimensional space.
 - Balanced index structures in multiple dimensions are challenging



 $\begin{array}{l} \langle 10,11\rangle \text{, } \langle 20,11\rangle \\ \langle 70,12\rangle \text{, } \langle 80,10\rangle \end{array}$

The R-tree

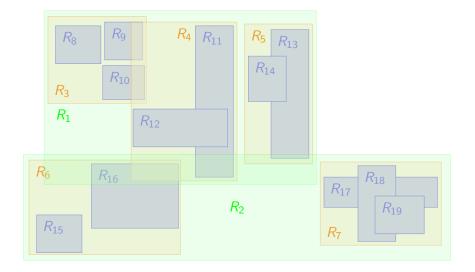
- The *R-tree* is a *tree-structured* index that remains *balanced* on *insertions* and *deletions*
- Each key stored in a leaf entry is intuitively a box, or collection of intervals, with one interval per dimension



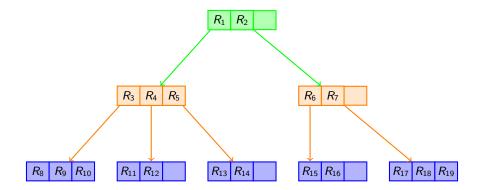
R-tree properties

- Leaf entry format: (n-dimensional bounding box, pointer to record)
 - Bounding box is the tightest bounding box for a data object
- Non-leaf entry format: (n-dim box, pointer to child node)
 - The box covers all boxes in child node (in fact, subtree)
- All leaves at same distance from root
- *Nodes* can be kept *50% full* (except root)
 - ► Can choose some parameter m that is ≤ 50%, and ensure that every node is at least m% full

R-tree example



R-tree example (cont.)



Search for objects overlapping box Q

Start at root

```
If current node is non-leaf
```

For each entry $\langle E, ptr \rangle$, if box E overlaps Q, search subtree identified by ptr

```
If current node is leaf
```

```
For each entry \langle E, rid \rangle, if E overlaps Q, rid identifies an object that might overlap Q
```

Note

May have to *search several subtrees* at each node! (In *contrast*, a B+tree equality search goes to *just one leaf*.)

```
Insert entry \langle B, ptr \rangle
```

Start at root and go down to "best-fit" leaf L

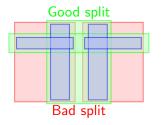
Go to *child* whose *box* needs *least enlargement* to *cover B*; *resolve ties* by going to *smallest area child*

If best-fit leaf L has space, insert entry and stop. Otherwise, split L into L_1 and L_2

Adjust entry for L in its parent so that the box now covers (only) L_1 Add an entry (in the parent node of L) for L_2 . (This could cause the parent node to recursively split.)

Splitting a node

- The *entries* in *node L* plus the *newly inserted* entry must be *distributed* between *L*₁ and *L*₂
- *Goal* is to *reduce likelihood* of *both* L₁ *and* L₂ being *searched* on subsequent queries
- *Redistribute* so as to *minimize area* of L_1 *plus* area of L_2



Redistribution

Exhaustive algorithm is *too slow*; *quadratic* and *linear heuristics* are used in practice

Comments on R-trees

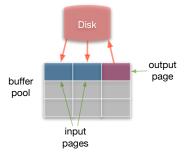
- Deletion consists of searching for the entry to be deleted, removing it, and if the node becomes under-full, deleting the node and then re-inserting the remaining entries
- Overall, works quite well for 2- and 3-D datasets
- Several variants (notably, R+ and R* trees) have been proposed; widely used
- Can *improve search performance* by using a *convex polygon* to *approximate query shape* (*instead* of a *bounding* box) and testing for *polygon-box intersection*.

Overview

- Sorting is probably the most classic problem in CS
 - Simple idea: impose a total order on a set of values
- It is a *classic problem* in *databases* too
 - Remember ISAM? First step is to sort the file
 - In fact, if you're bulk loading a B+tree, you're better off sorting the file first
- Useful as well for duplicate elimination
- Useful for *join evaluation* (*sort-merge* algorithm)
- But what if I have a *1GB relation* and *1MB of physical memory*?
 - Remember, its all about *minimising I/O*
 - (Or, why your algorithms class didn't tell you the whole truth)

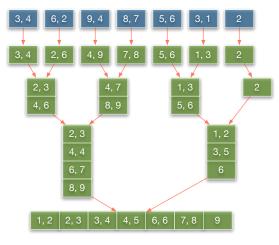
Two-way external merge sort

- Requires a *maximum of three* buffer pages and multiple passes over the data
- First pass: read one page, sort it, write it out
- Subsequent passes: read two pages, merge them, write out the result



How it works

- Each pass will read and write each page in the file
- *N* pages, so the number of passes is $\lceil \log_2 N \rceil + 1$
- So, the *total I/O cost* is $2N(\lceil \log_2 N \rceil + 1)$



But why only three pages?

- We have an *entire buffer pool* of *more than three pages*, can we utilise it?
 - Yes: N-way merge sort
- To sort a file of N pages using B buffer pool pages:
 - First pass: sorted runs of B pages each $\left(\begin{bmatrix} N \\ B \end{bmatrix} \right)$
 - Subsequent passes: merge B 1 runs (why?)

What is the I/O cost?

- Number of passes: $1 + \lceil \log_{B-1} \lceil \frac{N}{B} \rceil \rceil$
- $I/O \text{ cost: } 2N \cdot \text{ (Number of passes)}$
- For example: 108 pages in the file, 5 buffer pool pages
 - Pass 0: $\lceil \frac{108}{5} \rceil = 22$ sorted runs of 5 pages each
 - Pass 1: $\lceil \frac{22}{4} \rceil = 6$ sorted runs of 20 pages each
 - Pass 2: 2 sorted runs, 80 pages and 28 pages
 - Pass 3: final merge, done!

Sorting

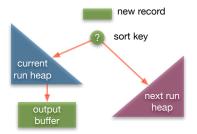
A bit of perspective

| | | 257 * 4,096 = 1,052,672 | | | | |
|---------------|-----|-------------------------|-----|------|-------|-------|
| Ν | B=3 | B=5 | B=9 | B=17 | B=129 | B=257 |
| 100 | 7 | 4 | 3 | 2 | 1 | 1 |
| 1,000 | 10 | 5 | 4 | 3 | 2 | 2 |
| 10,000 | 13 | 7 | 5 | 4 | 2 | 2 |
| 100,000 | 17 | 9 | 6 | 5 | 3 | 3 |
| 1,000,000 | 20 | 9 | 7 | 5 | 3 | 3 |
| 10,000,000 | 23 | 12 | 8 | 6 | 4 | 3 |
| 100,000,000 | 26 | 14 | 9 | 7 | 4 | 4 |
| 1,000,000,000 | 30 | 15 | 10 | 8 | 5 | 4 |

Are we done?

- No! We can actually do much better than this
- *Key observation*: we are using *main memory algorithm* (*e.g.*, quicksort) to sort pages in memory
 - But that doesn't minimise I/O, does it?
 - Wouldn't it be nice if we could generate sorted runs longer than memory?
 - ► Solution: heapsort (a.k.a. tournament or replacement sort)

How does heapsort work?



- Keep two heaps in memory, one for each run (the current and the next one)
- Sum of memory needed for the two heaps equals the buffer size
- *Keep adding* to the *current* run *until* we are *out of buffer space*
- When *buffer is full*, *swap* heaps and *iterate*

The algorithm

```
Initialisation: read B pages into the current heap
while (not finished) do {
 while (r = lowest key from current heap) {
  write r to the current run
  max = r
  get k from input
  if (k > \max) insert k into current heap
  else insert k into next heap
 }
 swap current and next heaps, max = 0
```

Sorting

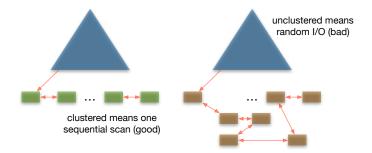
Heapsort observations

- What is the average length of a run?
 - Proven to be 2B (!)
- Quicksort is computationally cheaper
- But *heapsort* produces *longer runs* ٠
 - Minimises I/O
 - Remember, you should "forget" main memory methods when it comes to databases!

Good-old B+trees

- What if the *table to be sorted* has a *B*+*tree index on sort field*?
- Traverse the leaf pages and we're done!
 - ► Follow the *left-most pointers*, find the *low key*, *scan forward*
- Is this always a good idea?
 - ▶ If the *B*+tree is clustered, it's a great idea
 - Otherwise, it could lead to random I/O

Clustered vs. unclustered storage



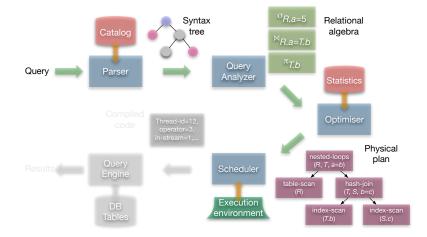
Summary of sorting

- Databases spend a lot of their time sorting
- In fact, they might *dedicate part of their buffer pool* for sorting data
 - Remember pinning buffer pool pages?
- External sorting minimises I/O cost
 - First you produce sorted runs, then you merge them
- The choice of internal sort matters as well
 - ► Yes, *quicksort* is *computationally cheap*
 - Though heapsort is computationally more expensive, it produces longer runs, which means less I/O
- Finally, *clustered B+trees* (when they exist) are a good way of *sorting in one sequential scan*

Overview

- A *physical plan* is what the *query engine* uses in order to *evaluate queries*
- In most cases, it is a tree of physical operators
 - Physical in the sense that they access and manipulate the raw, physical data
- Plenty of ways to formulate this tree
 - Identifying the "best" tree is the job of the query optimiser

Query cycle



Algebraic operators vs. physical operators

- A relational algebraic operator is a procedural abstraction of what should be retrieved
- The *physical operator* specifies how the retrieval will take place
- The same algebraic operator may map to multiple physical operators
- *Physical operators* for the *same algebraic operator* may be implemented using *different algorithms*
 - ► For instance: *join* → *physical join* → *sort-merge join*

Example

SQL query

| select | <pre>student.id, student.name</pre> |
|--------|---|
| from | student, course |
| where | <pre>student.cid = course.cid and</pre> |
| | course.name = 'ADBS' |

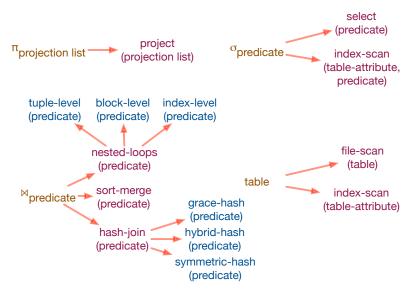
Algebraic expression

 π student.id,student.name (student $\bowtie_{student.cid=course.cid}$ $\sigma_{course.name='ADBS'}$ (course))

Algebraic operations

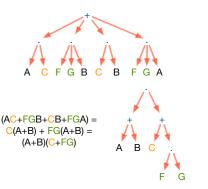
- $\pi_{student.id, student.name}$
- $\bowtie_{student.cid} = course.cid$
- $\sigma_{course.name} = `ADBS'$

Mappings to/of various physical operators



Math analogy

- Remember *factoring*?
- Same *arithmetic expression* can be *evaluated in different ways*
- If you map arithmetic expressions to *infix notation*, you have *different "plans"*



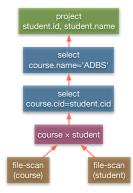
Physical plans

- Physical plans are trees of physical operators over the physical data
 - Just as arithmetic expressions are trees of arithmetic operators over numbers
- There are *different ways* of *organising trees* of *physical operators*
 - ► Just as there are *different ways* to *organise* a *mathematical expression*
- Physical plans are what produce query results

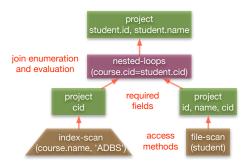
Here's a plan

SQL query

| select | <pre>student.id, student.name</pre> |
|--------|---|
| from | student, course |
| where | <pre>student.cid = course.cid and</pre> |
| | course.name = 'ADBS' |



Here's a better plan



SQL query

| select | <pre>student.id, student.name</pre> |
|--------|---|
| from | student, course |
| where | <pre>student.cid = course.cid and</pre> |
| | <pre>course.name = 'ADBS'</pre> |
| | |

Observations

- Certain *selection predicates* can be *incorporated* into the *access method*
- If a *field* is *not needed*, it is *thrown out* (why?)
- *More than two sources* need to be *combined* (even through a Cartesian product)
- The query plan includes operators not present in the original query
- Yes, the query specifies what should be retrieved
 - But how it is retrieved is an entirely different business

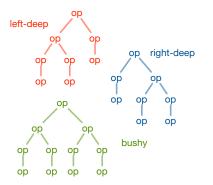
Issues

- Choice of order in which the physical operators are executed
 - Heuristics, access methods, optimisation
- Choice of algorithms whenever there are more than one
 - Again, optimisation (join enumeration, mainly)
- How are physical operators connected?
 - Different execution models
- What does a *connection* actually *imply*?
 - Pipelining (sometimes)
- What about *multiple readers* or even *concurrent updates* of the data?
 - Concurrency control (be patient ...)
- Finally, how is it all executed?
 - Query engine

A note on duplicates

- The *relational model* calls for *sets of tuples*
- The query language (SQL) does not
 - Remember "*distinct*"?
- Sets can be guaranteed on base relations by specifying key (integrity) constraints
- But what happens with *intermediate results*?
 - Set semantics are lost, intermediate results have bag semantics
 - But set semantics can always be imposed; they are just more expensive to ensure

Types of plan



- There are *two types of plan*, *according to their shape*
 - Deep (left or right)

Bushy

• Different shapes for different objectives

Plan objectives

- A *deep plan* is better for *pipelining*
 - Because, let's face it, it's a line!
- A *bushy plan* is better for *parallel computation*
 - Different branches can be executed concurrently
- But all of these *depend* on the *algorithms chosen*
 - And on the execution model

Summary

- A *plan* is what the *query engine* accepts as *input*
 - ... and what produces the query results
- The same algebraic expression can produce multiple plans
 - Because the same algebraic operator maps to multiple physical operators
- A physical operator implements an evaluation algorithm
- A *physical plan* does *not necessarily contain all* the *algebraic operators* of the query
 - ► More or fewer, depending on the available physical operators
- The optimiser chooses the best physical plan
- *Types* of plans are *classified* according to their *shape* and *evaluation objectives*

Overview

- Physical plans are trees of connected physical operators
- The execution model defines the interface of the connections
 - And how data is propagated from one operator to the next
- It also defines *how operators* are *scheduled* by the query engine
 - Different execution models map to different process execution paradigms

Execution models

Operator connections

- Operator *functionality*: relation in, relation out
- The *connections* are the *interface* through which the *input* is *read* and *propagated*
- In fact, there is a producer/consumer analogy



Pipelining

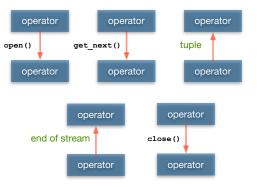
- *Pipelining* is the following process: *read*, *process*, *propagate*
- The opposite is to materialise intermediate results
- Pipelining *works in theory*, but *in practice* certain *intermediate relations* need to be *materialised*
 - ► This is called *blocking* (*e.g.*, sorting)
- The benefits of pipelining include
 - No buffering
 - ★ No intermediate relation is materialised
 - Faster evaluation
 - * Since nothing is materialised, no disk I/O
 - Better resource utilisation
 - ★ No disk I/O means more *in-memory operation*

What happens in practice

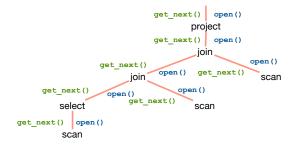
- *Pipelining* is *simulated* through the *operator interface*
- But different operations have different evaluation times
 - ► So there will be *some need for buffering*
- If we have *joins*, chances are the *plan will block*
 - ► We will see *why* that happens when talking about *join algorithms*

The iterator model

- Also known as a cursor
- Three basic calls
 - ▶ open()
 - > get_next()
 - ▶ close()
- Have you ever accessed a database through external code?
 - For example: exec sql declare cursor in embedded SQL, ResultSets in Java/JDBC, etc.



Call propagation



- All *calls* are propagated downstream
- The *query engine* makes the *calls* to the *topmost operator only*

Pure implementation

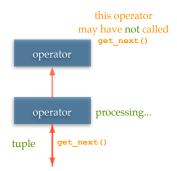
- The *iterator interface*, as described, is a *completely synchronous* interface
- A *pure implementation* means that all *operators reside in the same process space*
 - ► So *calls* can be *propagated downstream*
- But certain operators are "faster" than others
 - It could be the case that an asynchronous implementation could be more beneficial

Different implementations

- The iterator interface is what operators use to communicate
- But how it is implemented, can be entirely different
 - ► The *reason* is that there might be *need for buffering*
 - Three possibilities
 - * Push model (buffering in the operator making the calls)
 - * *Pull* model (buffering in the *operator accepting the calls*)
 - * *Streams* (buffering in the *connections*)

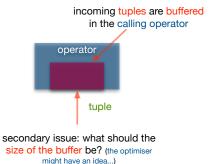
The push model

- Tuple *propagation begins* at the *lower levels* of the evaluation tree
- A *lower operator propagates* a tuple *as soon as it is done* with it
 - Does not "care" if the receiving operator has called get_next()



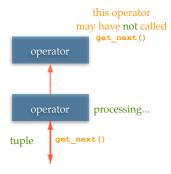
Buffering

• The main issue: *what happens* if the *lower operator* has *propagated* the tuple *before* the *operator above* it has called get_next()?



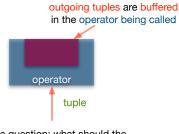
Stratis D. Viglas (University of Edinburgh)

The pull model



- The *inverse* of the *push model*
- If the *lower operator* is *done processing* a *tuple* it *does not propagate* it
 - It waits until the operator above it makes a get_next() call

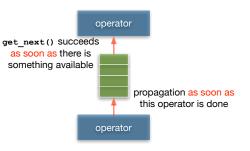
Buffering — again



same question: what should the size of the buffer be? (again, the optimiser might have an idea...) • The question this time: what happens if the lower operator is done processing the tuple before the operator above it calls get_next()?

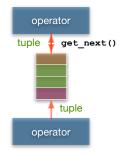
The stream model

- The *connections* become *first-class citizens*
- *Streams* are *queues* of *tuples* connecting the operators
- Propagations and get_next() calls are synchronised on each stream



Buffering — third time

- This time, there is no question!
- When the *lower operator* is *done*, it *propagates* the tuple
- When the *top operator* is *ready*, it calls get_next() on the incoming stream



Why all this?

- Pure iterator implementation
 - If an operator receives get_next() and is not ready, it blocks
 - ► In fact, the *entire plan blocks* (why?)
 - Assume there is a sort operation somewhere in the plan
 - ★ Congratulations, your plan is officially blocked
- Non-pure implementations
 - Operators act (almost) independently of one another
 - Depending on the implementation of the interface (push-, pull-, stream-based) there are different benefits
 - ★ There *could still be blocking*, but the *time during* which a *plan* is *blocked* is *minimised*
 - It could lead to a each operator running in its own process thread
 - ★ Though this is *not always* a *good idea*

Benefits of each model

Push model

- Minimises idle time of the operators (why?)
- Great for pipelining
- Pull model
 - Closest to a pure implementation
 - But still on-demand
- Streams model
 - Fully asynchronous to the operators, the synchronisation is on the streams
 - Highly parallelisable

Summary

- A physical plan is a tree of connected operators
- Operators need to communicate data to one another
- The *iterator interface* is the *means* of this *communication*
 - open(), close(), get_next()
- As with any *interface* there are *different ways of implementing* it, known as *execution models*
 - Push model
 - * Data propagated as soon as they are available
 - Pull model
 - * Data retrieved on demand
 - Stream model
 - * Asynchronous communication on the connections between operators

Overview

- The join operation is everywhere
 - Any single query with two or more sources will need to have a join (even in the form of a Cartesian product)
 - ► So common that certain DBMSs implement join indexes
- As a *consequence*, a DBMS spends *a lot of time* evaluating *joins*
- Probably the most optimised physical operator
- A physical operator can be mapped to different algorithms
- As is always the case, a good join algorithm minimises I/O
- *Choosing* a *join algorithm* is *not* as *straightforward*; the *choice* might *depend* on
 - ► The *cardinality* of the input, its *properties* (clustered, sorted, *etc.*) and any available *indexes*
 - Available memory

Overview (cont.)

- Choosing how to evaluate a single join is different than choosing the order in which joins should be evaluated
- The *query optimiser* spends *most of its time enumerating (ordering)* the *joins* in a query
 - In fact, the order in which joins are evaluated affects the choice of algorithm
 - The two are largely interconnected (more on that when discussing query optimisation)

Three classes of algorithms

- Iteration-based
 - Namely, nested loops join (in three flavours)
- Order-based
 - Sort-merge join (essentially, merging two sorted relations)
- Partition-based
 - Hash join (again, in three flavours)

Terminology

- We want to evaluate $R \bowtie S$, shorthand for R.a = S.b
 - Also known as an equi-join
- In algebra: $R \bowtie S = S \bowtie R$
 - Not true for the physical join: $cost(R \bowtie S) \neq cost(S \bowtie R)$
- Three factors to take into account
 - ► Input cardinality in tuples T_R and pages P_R
 - Selectivity factor of the predicate
 - * Think of it as the percentage of the Cartesian product propagated
 - Available memory

Nested loops join

- The *simplest way* to *evaluate a join*
- But it can still be optimised so that it minimises I/O
- Very useful for non-equi joins (the other two approaches will not work)
- Three variations
 - Tuple-level nested loops
 - Block-level nested loops
 - Index nested loops

It doesn't get simpler than this...

Tuple-level nested loops

for each tuple $r \in R$ do for each tuple $s \in S$ do if r.a == s.b then add $\langle r, s \rangle$ to the result

- *R* is the outer relation
- S is the inner relation

What is the cost?

- One scan over the outer relation
- For every tuple in the outer relation, one scan over the inner relation
- If relations are *not clustered*, then
 - $cost(R \bowtie S) = T_R + T_R \cdot T_S$
 - * Assume $T_R = 100,000, T_S = 50,000$, then cost = 5,000,100,000 I/Os
 - * At 10ms an I/O, that is 50,001,000 seconds, or, 14,000 hours

What about clustered storage?

- Much, much better; I/O is at a page level
- So, the *total cost* will be
 - $cost(R \bowtie S) = P_R + P_R \cdot P_S$
 - ▶ In the previous example, for 100 tuples per page, then $P_R = 1,000$, $P_S = 500$, cost = 501,000 I/Os
 - ► At 10ms an I/O, that is 5010 seconds, or about an hour and a half
- But we can *improve* that *even more*!
 - Block-level I/O and the buffer pool will work wonders

Here's an idea

- Assume we have *B pages available* in the buffer pool
- *Read as many outer relation pages as possible*; this constitutes a *block*
 - Put the pages of the block in the buffer pool, pin them
- Read the inner relation in pages
- Block size is B 2 pages (why?)
- Even more I/O savings

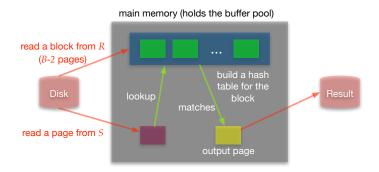
The Algorithm

Block-level nested loops

Assumption: B dedicated pages in the buffer pool, block size is B-2 pages

```
for each block of B - 2 pages of R do
for each page of S do {
for all matching in-memory tuples r \in R-block and s \in S-page
add \langle r, s \rangle to result
```

How it works



How much does it cost?

- The outer relation is still scanned once (P_R pages)
- The *inner relation* is scanned $\left\lceil \frac{P_R}{B-2} \right\rceil$ times
 - Each scan costs P_S I/Os
 - So, $cost(R \bowtie S) = P_R + P_S \cdot \left\lceil \frac{P_R}{B-2} \right\rceil$
 - Same example, $P_R = 1,000$, $P_S = 500$, assume a block size of 100 pages, then *number of I/Os is 6,500*
 - At 10ms per I/O, it will take 65 seconds

Key observation

- The *inner relation* is *scanned* a number of *times* that is *dependent on* the *size* of the *outer relation*
- So, the outer relation should be the smaller one
- Let's forget the ceilings and assume two relations: big and small
- Then we are comparing
 - $big + small \cdot \frac{big}{B-2}$ • $small + big \cdot \frac{small}{B-2}$
- And **big** > small
- *Remember*, $cost(R \bowtie S) \neq cost(S \bowtie R)$ when it comes to *physical* operators

Join algorithms

What if there is an index?

- Suppose the *inner relation* has an *index on the join attribute*
- We can use the index to evaluate the join
 - Remember, the *join predicate*, if we fix one of the join attribute values, is *just a selection*
- Scan the outer relation
 - Look at the join attribute's value and use it to perform an index lookup on the inner relation

The algorithm

```
Index nested loops

Assumption: there is an index on S.b

for each tuple r \in R do

for each tuple s \in S where r.a == s.b

add \langle r, s \rangle to the result
```

• Predicate evaluation is an *index lookup* in the *index* over *S*.*b*

What is the cost?

- Depending on whether the outer relation is clustered or not, P_R or $T_R I/Os$ to scan it
- Selectivity factor f: percentage of the Cartesian product propagated; this means that every outer tuple joins with $f \cdot T_S$ tuples
 - Depending on the index, each lookup will be, say, avg_lookup I/Os
- If *R* is *clustered*
 - $cost(R \bowtie S) = P_R + T_R \cdot f \cdot T_S \cdot avg_lookup$
- If *R* is not clustered
 - $cost(R \bowtie S) = T_R + T_R \cdot f \cdot T_S \cdot avg_lookup$

Index nested loops

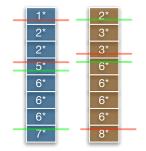
- If the *selectivity factor* and the *average lookup cost* are *small*, then the *cost* is *essentially a (few) scan(s)* of the *outer relation*
- If the *outer relation* is the *smaller one*, it leads to *significant I/O savings*
- Again, it is the *job* of the *query optimiser* to *figure out if this is the case*

Sort-merge join

- Really *simple idea*
- The join is evaluated in two phases
 - ► *First*, the two *input relations* are *sorted* on the *join attribute*
 - Then, they are merged and join results are propagated
- External sorting can be used to sort the input relations
- The *merging phase* is a *straightforward generalisation* of the *merging phase* used in *merge-sort*

How it works

- Key idea: there *exist groups* in the *sorted relations* with the *same value* for the *join attribute*
- We need to *take that* into *account* when *merging*
 - The reason is that we will have to do some backtracking when generating the complete result



The algorithm

Merge-join

```
r \in R, s \in S, gs \in S
while (more tuples in inputs) do {
 while (r.a < gs.b) do advance r
 while (r.a > gs.b) do advance gs //a group might begin here
 while (r.a == gs.b) do {
  s = gs // mark group beginning
  while (r.a == s.b) do // while in group
      add \langle r, s \rangle to the result; advance s // produce result
  advance r // move forward
 gs = s // candidate to begin next group
```

What is the cost?

- We know the cost of externally sorting either relation: 2 · P_R · log P_R, or 2 · P_S · log P_S
- The *merge phase* is essentially *one scan* of *each sorted input*: *P*_R or *P*_S (these scans are always clustered)
- $cost(R \bowtie S) = P_R \cdot (2 \cdot \log P_R + 1) + P_S \cdot (2 \cdot \log P_S + 1)$
 - ▶ Running example: $P_R = 1,000$, $P_S = 500$, 100 buffer pool pages to sort, the *number of I/Os is 7,500*
 - At 10ms an I/O, this is one minute and fifteen seconds (about the same as block nested loops)

A few issues

- If there are *large groups* in the *two relations*, then we *may* have to *do a lot of backtracking*
 - Performance will suffer due to possible extra I/O
 - Hopefully, pages will be in the buffer pool
- Most relations can be sorted in 2-3 passes
 - Which means that we can compute the join in 4 passes max (almost regardless of input size!)
 - In fact, we can combine the final merge of external sorting with the merging phase of the join and save even more I/Os

Hash join

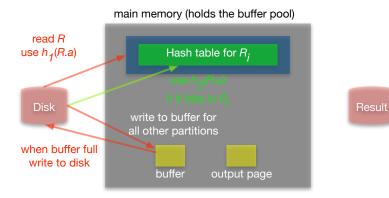
- Partition-based join algorithms
- Key idea: *partition R and S* into *m partitions*, *R_i* and *S_i*, so that every *R_i* fits in memory
 - Observation: joining tuples will fall into the same partition
- Then, for every R_i load it in memory, scan S_i and produce the join results
- Three flavours: Simple hash join, grace hash join, hybrid hash join

The simple algorithm

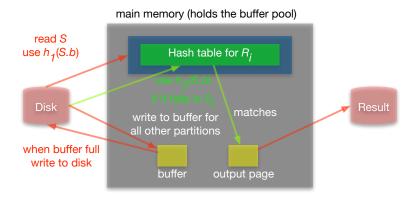
Simple hash join

Assumption: m partitions, each partition P_i fits in main memory for all partitions P_i , $i \in [1, m]$ for each $r \in R$ read r and apply hash function $h_1(r.a)$ if r falls into P_i apply hash function $h_2(r.a)$ and put it in an in-memory hash table for P_i otherwise, write it back out to disk for each $s \in S$ read s and apply hash function $h_1(s.b)$ if s falls into P_i apply hash function $h_2(s.b)$ and for all matching tuples $r \in P_i$, add $\langle r, s \rangle$ to the result otherwise, write it back out to disk

How it works — partitioning R, iteration i



How it works — paritioning and joining S, iteration i



What is the cost?

- Assume equal partition sizes, input T, P_T pages
- For *m partitions*, we will make *m passes* over each input
 - For the first pass:

* Read P_T pages, write $P_T - \frac{P_T}{m}$ pages: $2P_T - \frac{P_T}{m}$ I/Os

For the second pass:

* Read $P_T - \frac{P_T}{m}$, write $P_T - \frac{P_T}{m} + P_T - 2\frac{P_T}{m}$ pages: $2P_T - 3\frac{P_T}{m}$ I/Os

- Pass *i*: $2P_T (2i 1) \frac{P_T}{m}$ I/Os
- In the end, $m(m+1)P_T$ I/Os
- For two relations R and S, total cost is $m(m+1)(P_R+P_S)$
- Makes sense if *m* is small, or we have a lot of memory
- Effectively, this is nested loops join
 - But the number of iterations is decided by the number of partitions, not the input sizes!

The "grace" algorithm

Grace hash join

```
for each r \in R read r and add it to the buffer page for h_1(r.a)
for each s \in S read s and add it to the buffer page for h_1(s.b)
for i = 1, ..., m do {
for each r \in R_i read r and insert it into a hash table using h_2(r.a)
for each s \in S_i do {
read s, probe the hash table using h_2(s.b)
for all matching tuples r \in R_i add \langle r, s \rangle to the result
}
clear hash table
```

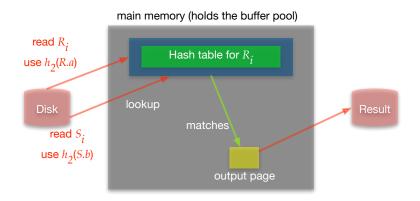
How it works — partitioning R

main memory (holds the buffer pool) read Ruse $h_1(R.a)$ Disk R_1 R_2 ... R_m as soon as a page for R_i fills up write it to disk

How it works — partitioning S

read S use $h_1(S.b)$ Disk S1 S2 ... Sm as soon as a page for S_i fills up write it to disk

How it works — joining



What is the cost?

- Scan R and write it to disk, so $2 \cdot P_R$
- Do the same for S, so $2 \cdot P_S$
- Read R in partition-by-partition, so P_R
- Scan S partition-by-partition and probe for matches, so P_S
- $cost(R \bowtie S) = 3 \cdot (P_R + P_S)$
 - Same example, $P_R = 1,000$, $P_S = 500$, cost is 4,500 I/Os
 - At 10ms an I/O the join will take 45 seconds to evaluate

Memory requirements

- Objective: the hash table for a partition must fit in memory
 - Minimise partition size by maximising number of partitions
- What are the optimum sizes?
 - For *B* buffer pool pages, maximum number of partitions m = B − 1 (why?)
- Size of each partition is $\left\lceil \frac{P_R}{B-1} \right\rceil$
- Size of the hash table is $\left[\frac{f \cdot P_R}{B-1}\right]$ ($f = fudge \ factor$ to capture the increase in partition size due to the hash table)
- During the *probing phase*, in addition to the hash table, we need *one* page to read S, plus one page for output

• So,
$$B > \left\lceil \frac{f \cdot P_R}{B-1} \right\rceil + 2 \Rightarrow B > \sqrt{f \cdot P_R}$$

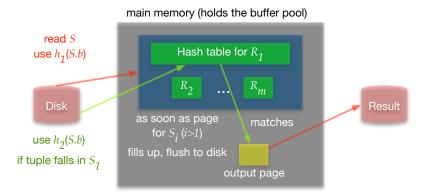
Hybrid hash join

- An improvement over hash join if there is extra memory
 - Minimum amount of memory for hash join $B > \sqrt{f \cdot P_R}$
 - Suppose that $B > \frac{f \cdot P_R}{k}$, for some integer k
 - Divide R into k partitions of size $\frac{P_R}{k}$ (k + 1 buffer pool pages needed)
 - This leaves B (k + 1) extra buffer pool pages

How it works

- Suppose that $B (k+1) > \frac{f \cdot P_R}{k}$
 - ► We have enough memory during partitioning to hold an in-memory hash table of size B (k + 1) pages
- Idea: keep R₁ in memory at all times
- While partitioning S, if a tuple falls into S_1 , don't write it to disk; instead probe the hash table for R_1 for matches
- For all partitions $R_i, S_i, i > 2$, continue as in hash join

How it works — partitioning and joining



Savings over grace hash join

- Essentially, *reduces* the *number* of *full passes*
- Running example, $P_R = 1,000$, $P_S = 500$, assume 300 pages in the buffer pool
- Choose the smaller relation, S
- Two partitions for it, each 250 pages
 - ▶ But one will stay in memory; so, cost is 500+250=750 I/Os
- Scan R, use two partitions, each 500 pages
 - ▶ But the *first one* is *not written* to disk; so cost is 1,000+500=1500 I/Os
- Join the two on-disk partitions, cost 250+500=750 I/Os
- Total cost 750+1500+750=300 I/Os
- At 10ms an I/O, this is half a minute

On predicates

• The algorithms we talked about will work on equi-join predicates

- If there are no equi-join predicates (inequality joins) the only algorithm that will work is nested loops (why?)
- If there are indexes on the inequality join predicate's attributes, we can use index nested loops and revert the join to multiple scans
 - * Hoping that we will have buffer pool hits
 - ★ Remember access patterns and page replacement policy?
- Luckily, in a typical query workload there will mostly be equi-join predicates

On pipelining

- *Pipelining* is *great*, but it *cannot always be achieved*
- All three algorithms will essentially block at some point
 - ► In the *best case*, *between matches*
 - ► In the *worst case*, *until after a few scans* of the input relations
- This is *not necessarily bad*; in fact, *even* if the *algorithms block*, the *time needed* to compute the complete join result *might be less*
- In reality, more than two stages of pipelining can rarely be obtained in a single plan

Summary

- The physical join is the most optimised physical evaluation operator
 - Because a DBMS spends most of its time evaluating joins
- Three main classes of algorithms
 - Iteration-based, order-based, partition-based
- Three main *choice criteria*
 - Physical layout, indexes, available memory

Summary (cont.)

• Iteration-based methods

- Essentially, nested loops
- Very simple to implement, but if implemented poorly very inefficient
- ► But also *very useful* because they *evaluate non-equi-join predicates*

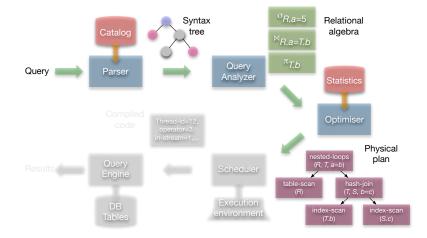
• Order-based methods

- Sort the inputs, merge them afterwards
- Well-behaved cost 3-4 passes over the data will do the trick

Summary (cont.)

- Partition-based methods
 - Simple hash join, Grace hash join, and hybrid hash join
 - ► If there is extra memory, hybrid hash join's behaviour is excellent
- *Figuring out* the *best join algorithm* for a *particular pair* of inputs is the *job* of the *query optimiser*
- Which, along with *good implementations*, will *choose the one* that *evaluates a join in 30 seconds and not in 14,000 hours*

Query cycle



Query optimiser

- The *query optimiser* is the *heart* of the *evaluation engine*
 - Yes, the physical operators get the job done
 - ▶ Yes, the *execution model* makes sure the *operators* actually *run*
 - ▶ But, unless the *query optimiser decides on those things*, the query will never run
 - And the *decision* needs to be a *good one*

Decisions

- Two crucial decisions the optimiser makes
 - The order in which the physical operators are applied on the inputs (*i.e.*, the plan employed)
 - The algorithms that implement the physical operators
- These *two decisions* are *not independent*
 - ▶ In fact, one affects the other in more ways than one

Cost-based query optimisation

- The paradigm employed is cost-based query optimisation
 - Simply put: *enumerate* alternative *plans*, *estimate* the *cost* of *each plan*, *pick* the *plan* with the *minimum cost*
- For *cost-based optimisation*, we need a *cost model*
 - Since what "hurts" performance is I/O, the cost model should use I/O as its basis
 - Hence, the cardinality-based cost model
 - * Cardinality is the number of tuples in a relation

Overview

Plan enumeration

- Plan enumeration consists of two parts (again, not necessarily *independent* from one another)
 - Access method selection (i.e., what is the best way to access a relation that appears in the query?)
 - Join enumeration (i.e., what is the best algorithm to join two relations, and when should we apply it?)
- Access methods, join algorithms and their various combinations define a search space
 - The search space can be huge
 - Plan enumeration is the exploration of this search space

Search space exploration

• As was stated, the *search space* is *huge*

- Exhaustive exploration is out of the question
- Because it could be the case that exploring the search space might take *longer than* actually *evaluating* the query
- The way in which we explore the search space describes a query optimisation method
 - * Dynamic programming, rule-based optimisation, randomised exploration, ...

Just an idea ...

- A query over *five relations*, only *one access method*, only *one join algorithm*, only *left-deep plans*
 - Remember, $cost(R \bowtie S) \neq cost(S \bowtie R)$
 - So, the number of *possible plans* is 5! = 120
 - ► If we add one extra access method, the number of possible plans becomes 2⁵ · 5! = 3840
 - ► If we add one *extra join algorithm*, the number of *possible plans* becomes 2⁴ · 2⁵ · 5! = 61440

Cardinality-based cost model

- A *cardinality-based cost model* means we need *good ways of* doing the following
 - Using cardinalities to estimate costs (e.g., accurate cost functions)
 - Estimating output cardinalities after we apply certain operations (e.g., after a selection the cardinality will change; it will not change after a projection)
 - ★ Because these output cardinalities will be used as inputs to the cost functions of other operations

Overview

Cardinality estimation

- An entire area of query optimisation
- Largely a *matter of statistics*
- It has triggered the "percentage wars"
 - "This estimation technique is within x% of the true value with a y%probability"
- Fact: the better the statistics, the better the decisions
- Another fact: errors in statistics propagate exponentially; after 4 or 5 joins, you might as well flip a coin
- Third fact: cost functions are discontinuous, so in certain scenarios only perfect statistics will help

Are we done?

- The previous issues were only a subset of the problems an optimiser solves
 - We also need to worry about certain properties of the data
 - ★ For instance, if we use a B+tree as an access method, then we won't have to sort (e.g., interesting orders in System R)
 - * If we use a *hash join later on* the *order is spoiled*
 - ★ So we will *have to sort again*
 - Depending on the algorithm and the environment, we need to allocate memory
- And as if all these were not enough, optimisation time assumptions do not necessarily hold at run time

The final nail ...

- These are *all* for *one query*
- Now, imagine a system doing that for 1000 queries
 - Simultaneously
- And it *all* has to be *done fast*
 - Once a decision is made, it cannot be undone

Conclusion

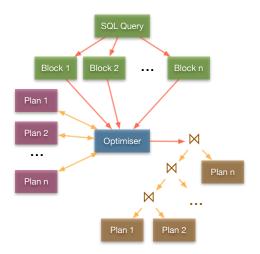
- Query optimisation is a very, very hard problem
- But *without it* a DBMS is doomed to seriously sub-optimal performance
- The problem is not nearly solved
 - All we have is decent optimisation strategies
 - And decent sub-problem solutions
- Fact: rarely will an optimiser pick the "best" plan
 - But it will almost always pick a plan with good performance and stay away from bad choices
 - At the end of the day, thats what counts

The agenda

- Mapping SQL queries to relational algebra
 - Query blocks, uncorrelated vs. correlated queries
- Optimisation of a single query block
- Equivalence rules
- Statistics and cardinality estimation
- Search space exploration
 - Dynamic programming (System-R)

SQL decomposition

- *SQL queries* are *optimised* by *decomposing* them into a *collection* of *query blocks*
- A *block* is *optimised* in *isolation*, *resulting* in a *plan* for a *block*
- *Plans* for *blocks* are *combined* to form the *complete plan* for the query



What is a block?

- An SQL query with no nesting
- Exactly one select-clause
- Exactly one from-clause
- At most one
 - Where-clause in conjunctive normal form
 - Group by-/sort by-clause
 - Having-clause

Example

Sample schema

- Sailors (sid, sname, rating, age)
- Boats (bid, bname, color
- Reserves (sid, bid, day, rname)

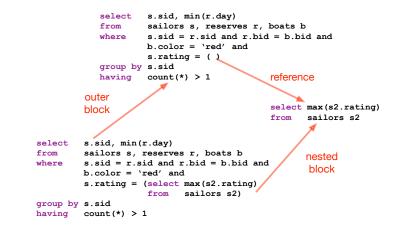
Example

For each sailor with the highest rating over all sailors, and at least two reservations for red boats, find the sailor id and the earliest date on which the sailor has a reservation for a red boat.

SQL query

| select | s.sid, min(r.day) |
|----------|--|
| from | sailors s, reserves r, boats b |
| where | s.sid = r.sid and r.bid = b.bid and |
| | b.color = 'red' and |
| | <pre>s.rating = (select max(s2.rating) from sailors s2)</pre> |
| group by | s.sid |
| having | count(*) > 1 |
| | |

Two blocks in the query



Single block optimisation — step 1

SQL query

| select | s.sid, min(r.day) |
|----------|--|
| from | sailors s, reserves r, boats b |
| where | s.sid = r.sid and r.bid = b.bid and |
| | b.color = 'red' and |
| | <pre>s.rating = (select max(s2.rating) from sailors s2)</pre> |
| group by | s.sid |
| having | count(*) > 1 |

Relational algebra

```
\pi_{s.sid,\min(r.day)}(
```

```
having<sub>count(*)>2</sub>(
```

```
group by<sub>s.sid</sub>(
```

```
\sigma_{s.sid=r.sid\wedge r.bid=b.bid\wedge b.color=red\wedge s.rating=nested-value}
```

sailors × *reserves* × *boats*))))

- *Express* the query in relational algebra
- More specifically, extended relational algebra

Single block optimisation — step 2

Relational algebra — before

 $\pi_{s.sid,\min(r.day)}($ $having_{count(*)>2}($ $group by_{s.sid}($ $\sigma_{s.sid=r.sid\land r.bid=b.bid\land b.color=red\land s.rating=nested-value}($ $sailors \times reserves \times boats))))$

Relational algebra — after

 $\pi_{s.sid}($

 $\sigma_{s.sid=r.sid \land r.bid=b.bid \land b.color=red \land s.rating=nested-value}$

 $sailors \times reserves \times boats))$

- *Ignore* the aggregate operations
 - They only have meaning for the complete result
 - Convert the query into a subset of relational algebra called σπ×

Single block optimisation — step 3

- Use *equivalence rules* to identify *alternative ways* of *formulating the query*
- "Plug in" algorithms
- Enumerate plans
- Estimate the cost of each plan
- *Pick* the *one* with the *minimum cost*

Equivalence rules

- Essentially, every query block consists of three things
 - Cartesian product of all relations in the from-clause
 - Selection predicates of the where-clause
 - Projections of the select-clause
- The *equivalence rules define* the *space* of *alternative plans* considered by an optimiser
 - ► In other words, the *search space of a query*

Selection and projections

• Cascading of selections

•
$$\sigma_{c_1 \wedge c_2 \wedge \ldots \wedge c_n}$$
 (R) $\equiv \sigma_{c_1} (\sigma_{c_2} (\ldots (\sigma_{c_n} (R))))$

• Commutativity

$$\blacktriangleright \sigma_{c_1} (\sigma_{c_2} (R)) \equiv \sigma_{c_2} (\sigma_{c_1} (R))$$

• Cascading of projections

$$\blacktriangleright \pi_{a_1} (R) \equiv \pi_{a_1} (\pi_{a_2} (\dots (\pi_{a_n} (R)) \dots)$$

• iff
$$a_i \subseteq a_{i+1}$$
, $i = 1, 2, \ldots n-1$

Cartesian products and joins

Commutativity

- $\blacktriangleright R \times S \equiv S \times R$
- $R \bowtie S \equiv S \bowtie R$
- Assosiativity
 - $R \times (S \times T) \equiv (R \times S) \times T$
 - $R \bowtie (S \bowtie T) \equiv (R \bowtie S) \bowtie T$
- Their combination

►
$$R \bowtie (S \bowtie T) \equiv R \bowtie (T \bowtie S) \equiv (R \bowtie T) \bowtie S$$

 $\equiv (T \bowtie R) \bowtie S$

Among operations

- Selection-projection commutativity
 - π_a (σ_c (R)) $\equiv \sigma_c$ (π_a (R))
 - ▶ iff every attribute in c is included in the set of attributes a
- Combination (join definition)
 - $\sigma_c (R \times S) \equiv R \bowtie_c S$
- Selection-Cartesian/join commutativity
 - $\sigma_c (R \times S) \equiv \sigma_c(R) \bowtie S$
 - ▶ iff the *attributes in c* appear *only in R* and *not in S*
- Selection distribution/replacement
 - ► $\sigma_c(R \bowtie S) \equiv \sigma_{c_1 \land c_2} (R \bowtie S) \equiv \sigma_{c_1} (\sigma_{c_2} (R \bowtie S)) \equiv \sigma_{c_1}(R) \bowtie \sigma_{c_2}(S)$
 - ▶ iff c₁ is relevant only to R and c₂ is relevant only to S

Among operations (cont.)

- Projection-Cartesian product commutativity
 - π_a ($R \times S$) $\equiv \pi_{a_1}(R) \times \pi_{a_2}(R)$
 - ▶ iff a₁ is the subset of attributes in a appearing in R and a₂ is the subset of attributes in a appearing in S
- Projection-join commutativity
 - π_a ($R \bowtie_c S$) $\equiv \pi_{a_1}(R) \bowtie_c \pi_{a_2}(R)$
 - ▶ iff same as before and every attribute in c appears in a
- Attribute elimination
 - $= \pi_a(R \bowtie_c S) \equiv \pi_a(\pi_{a_1}(R) \bowtie_c \pi_{a_2}(S))$
 - ▶ iff a₁ subset of attributes in R appearing in either a or c and a₂ is the subset of attributes in S appearing in either a or c

What do we have and what do we need?

We have

- ► A way to *decompose SQL queries* into *multiple query blocks*
- ► A way to *map a block* to *relational algebra*
- Equivalence rules between different algebraic expressions, i.e., a search space
- We need
 - ► A way to *estimate the cost* of *each alternative* expression
 - * *Depending* on the *algorithms* used
 - A way to *explore* the *search space*

Cost estimation

- A *plan* is a tree of operators
- *Two parts* to *estimating* the *cost* of a plan
 - For each node, estimate the cost of performing the corresponding operation
 - For each node, estimate the size of the result and any properties it might have (e.g., sorted)
- Combine the estimates and produce an estimate for the entire plan



Cost and cardinality

- We have seen various storage methods and algorithms
 - And know the cost of using each one, depending on the input cardinality
- The problem is estimating the output cardinality of the operations
 - Namely, selections and joins

Selectivity factor

- The *maximum number of tuples* in the *result* of any *query* is the *product* of the *cardinalities* of the *participating relations*
- Every *predicate* in the *where-clause* eliminates some of these *potential results*
- Selectivity factor of a single predicate is the ratio of the expected result size to the maximum result size
- Total result size is estimated as the maximum size times the product of the selectivity factors
- Key assumption: the predicates are statistically independent

How it works

SQL queryselect $a_1, a_2, \dots a_k$ from $R_1, R_2, \dots R_n$ where P_1 and P_2 and \dots and P_m

Maximum output cardinality $|R_1| \cdot |R_2| \cdot \ldots \cdot |R_n|$

Selectivity factor product $f_{P_1} \cdot f_{P_2} \cdot \ldots \cdot f_{P_m}$

Estimated output cardinality $(f_{P_1} \cdot f_{P_2} \cdot \ldots \cdot f_{P_m}) \cdot |R_1| \cdot |R_2| \cdot \ldots \cdot |R_n|$

Various selectivity factors

- column = value $\rightarrow \frac{1}{\#keys(column)}$
 - Assumes uniform distribution in the values
 - Is itself an approximation
- $column_1 = column_2 \rightarrow \frac{1}{\max(\#keys(column_1), \#keys(column_2)))}$
 - Each value in column₁ has a matching value in column₂; given a value in column₁, the predicate is just a selection
 - Again, an *approximation*

Various selectivity factors (cont.)

- column > value → (high(column)-value) (high(column)-low(column))
- $value_1 < column < value_2 \rightarrow \frac{(value_2 value_1)}{(high(column) low(column))}$
- column in list \rightarrow number of items in list times s.f. of column = value
- column in sub-query → ratio of subquery's estimated size to the number of keys in column
- not (predicate) \rightarrow 1 (s.f. of predicate)
- $P_1 \vee P_2 \to f_{P_1} + f_{P_2} f_{P_1} \cdot f_{P_2}$

Key assumptions made

- The values across columns are uncorrelated
- The values in a single column follow a uniform distribution
- Both of these assumptions rarely hold
- The first assumption is hard to lift
 - Only recently have researchers started tackling the problem
- The *uniform distribution* assumption can be *lifted* with *better statistical methods*
 - ► In our case, *histograms*

What we would like

Estimated distribution (uniform assumption)

9.00 6.75 4.50 2.25 0 0 2 3 4 5 6 7 8 9 13 14 T 10 11 12

True distribution

Cardinality estimation

Lifting the uniform distribution assumption

- At the *basic level*, all we *need* is a *collection* of *(value,* frequency) pairs
- Which is just a relation!
 - So, scan the input and build it
- But this is *unacceptable*
 - Because the size might be comparable to the size of the relation
 - And we need to answer queries about the value distribution fast

parts

| name | color | stock |
|------|-------|-------|
| bolt | red | 10 |
| bolt | green | 5 |
| nut | blue | 4 |
| nut | black | 10 |
| nut | red | 5 |
| nut | green | 10 |
| cam | blue | 5 |
| cam | green | 10 |
| cam | black | 10 |

parts.color

| value | freq |
|-------|------|
| red | 2 |
| green | 3 |
| blue | 2 |
| black | 2 |

parts.stock

| value | freq |
|-------|------|
| 10 | 4 |
| 5 | 3 |
| 4 | 1 |

Histograms

- Elegant data structures to capture value distributions
 - Not affected by the uniform distribution assumption (though this is not entirely true)
- They offer trade-offs between size and accuracy
 - ► The *more memory* that is dedicated to a histogram, the *more accurate* it is
 - But also, the more expensive to manipulate
- Two basic classes: equi-width and equi-depth

Desirable histogram properties

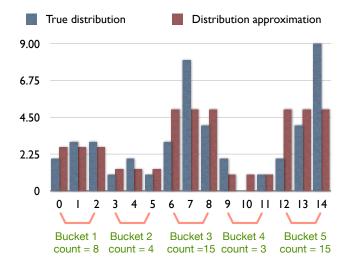
• Small

- Typically, a DBMS will allocate a single page for a histogram!
- Accurate
 - Typically, less than 5% error
- Fast access
 - Single lookup access and simple algorithms

Mathematical properties

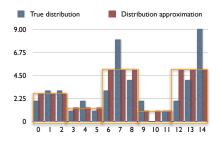
- A histogram approximates the value distribution for attribute X of table T
- The *value distribution* is *partitioned* into a number of *b subsets*, called *buckets*
- There is a *partitioning constraint* that *identifies how* the *partitioning* takes place
 - Different constraints, lead to different classes of histograms
- The *values* and *frequencies* in *each bucket* are *approximated* in some *common fashion*

Equi-width histogram



Equi-width histogram construction

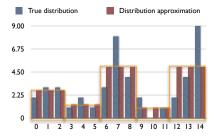
- The *total range* is *divided* into *sub-ranges* of *equal width*
- Each *sub-range* becomes a *bucket*
- The total number of tuples in each bucket is stored



| min | max | count |
|-----|-----|-------|
| 0 | 2 | 8 |
| 3 | 5 | 4 |
| 6 | 8 | 15 |
| 9 | 11 | 3 |
| 12 | 14 | 15 |

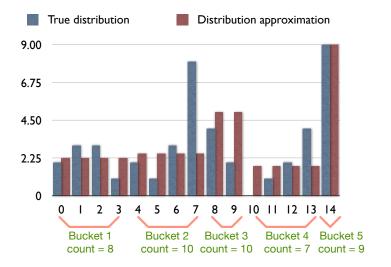
Equi-width histogram estimation

- To *estimate* the output cardinality of a range query
 - The starting bucket is identified
 - The histogram is then scanned forward until the ending bucket is identified
 - The numbers of tuples in the buckets of the range are summed
 - Within each bucket the uniform distribution assumption is made
- $6 \le v \le 10$: $\frac{3}{3} \cdot 15 + \frac{2}{3} \cdot 3 = 17$



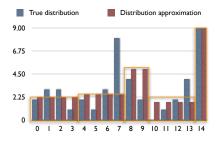
| min | max | count |
|-----|-----|-------|
| 0 | 2 | 8 |
| 3 | 5 | 4 |
| 6 | 8 | 15 |
| 9 | 11 | 3 |
| 12 | 14 | 15 |

Equi-depth histogram



Equi-depth histogram construction and estimation

- The *total range* is *divided* into *sub-ranges* so that the *number of tuples* in *each range* is (approximately) *equal*
- Each *sub-range* becomes a *bucket*
- The *same schema* as in *equi-width* histograms is used
- In fact, the *same algorithm* is used for *estimation* (!)
- $6 \le v \le 10$: $\frac{2}{4} \cdot 10 + \frac{2}{2} \cdot 10 + \frac{1}{4} \cdot 7 \approx 17$



| min | max | count |
|-----|-----|-------|
| 0 | 3 | 8 |
| 4 | 7 | 10 |
| 8 | 9 | 10 |
| 10 | 13 | 7 |
| 14 | 14 | 9 |

Comparison

- Equi-depth histograms are generally better than equi-width
 - Buckets with frequently occurring values contain fewer values
 - Infrequently occurring values are approximated less accurately (but the error is less significant)
 - So the uniform distribution assumption within each bucket leads to better approximation

What do we have and what do we need?

• We have

- A way to decompose a query
- A way to *identify* equivalent, *alternative representations* of it (*i.e.*, a search space)
- A statistical framework to estimate cardinalities
- ► A *cost model* to *estimate* the *cost* of an alternative
- We need
 - A way to *explore* the *search space*
 - Dynamic programming

Dynamic programming

- In the beginning, there was System R, which had an optimiser
- System R's optimiser was using dynamic programming
 - An efficient way of exploring the search space
- *Heuristics*: use the *equivalence rules* to *push down selections* and *projections*, *delay Cartesian products*
 - Minimise input cardinality to, and memory requirements of the joins
- Constraints: left-deep plans, nested loops and sort-merge join only
 - Left-deep plans took better advantage of pipelining
 - Hash-join had not been developed back then

Interesting orders

- If there is an *order by* or *group by* clause on an *attribute*, we say that this *attribute* has an *interesting order* associated with it
 - Interesting, because depending on the access method we can get away with fewer physical operations (e.g., sorting)
- The *same holds* for *attributes* participating in a *join*
 - Again, *interesting* because we can *use* the *access method* in *evaluating* the join

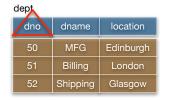
Dynamic programming steps

- *Identify* the *cheapest* way to *access every* single *relation* in the query, *applying local predicates*
 - For every relation, keep the cheapest access method overall and the cheapest access method for an interesting order
- For every access method, and for every join predicate, find the cheapest way to join in a second relation
 - ► For every join result keep the cheapest plan overall and the cheapest plan in an interesting order
- Join in the rest of the relations using the same principle

An example





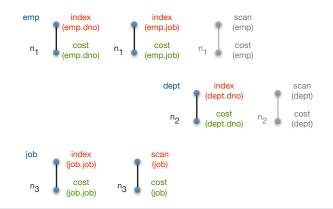


name, salary, job title, department name of employees who are clerks and work in departments in Edinburgh

local predicates



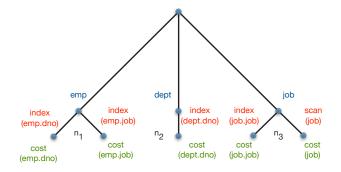
Access methods and local predicates



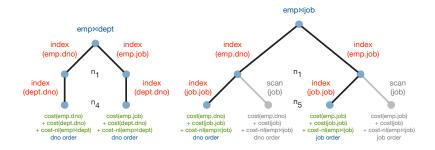
Scanning emp is the most expensive method for emp; emp.dno and emp.job are interesting orders

- Scanning dept is the most expensive method for dept; dept.dno is an interesting order
- Scanning job is the cheapest method for job; but, job.job is an interesting order

Search tree for access methods

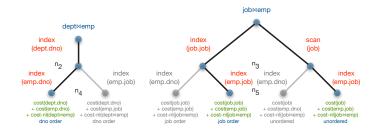


Join enumeration for relation *emp* (nested loops join)



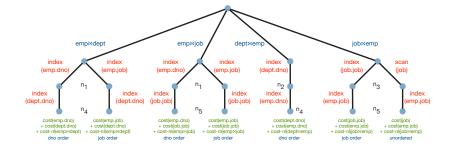
- Both emp \u2295 dept results are in different interesting orders so they are propagated
- Only the cheapest result in any interesting order is propagated for each pair of inputs

Join enumeration for relations *dept*, *job* (nested loops)

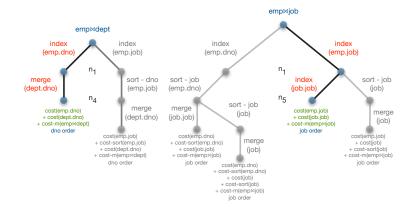


- ۰ $cost(emp \bowtie dept) \neq cost(dept \bowtie emp)$ so we will enumerate dept's joins even though we have an alternative for generating the same result (same for *job* \bowtie *emp*)
- ۰ Both dept \bowtie emp results in the same order, only one propagated
- ۲ Since there is no dept \bowtie job predicate in the query, that join is not enumerated (same for job \bowtie dept)
- ۲ The unordered result for job emp is propagated because it is the cheapest overall

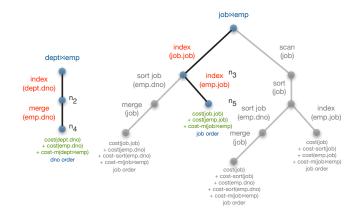
Search tree — 2 relations, nested loops join



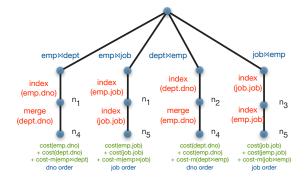
Join enumeration for relation *emp* (sort-merge)



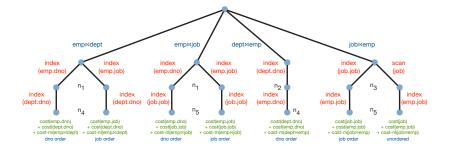
Join enumeration for relations *dept*, *job* (sort-merge)



Search tree — 2 relations, sort-merge join



Search tree — 2 relations, both join methods



For each pair or relations, for each different join order and for each interesting order for that pair one plan is propagated
 An unordered result is only propagated if it is the cheapest overall for a pair in a given join order

Three relations

• Repeat the process

- For every pair of two relations
- For every join method
- ► For every access method of the remaining relation
- Find the cheapest way to join the third relation with the pair
 - * Estimate cardinalities
 - ★ *Estimate* the *cost* of computing the *join*
- Keep the cheapest choice for every interesting order and the cheapest for the unordered case if it is the cheapest overall

Rule-based optimisation

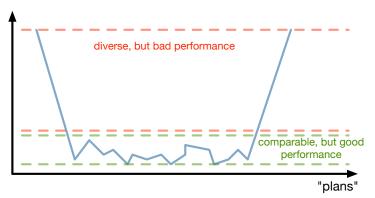
- Basically an issue of *if-then rules*
 - If (condition list) then apply some transformation to the plan constructed so far
 - *Estimate* the *cost* of the *new plan*, *keep* it *only if* it is *cheaper than* the *original*
 - ► The order in which the rules are applied is significant
 - As a consequence, rules are applied by precedence
 - * For instance, *pushing down selections* is given *high precedence*
 - Combining two relations with a Cartesian product is given low precedence

Randomised exploration

- Mostly useful in big queries (more than 15 joins or so)
- The *problem* is one of *exploring a bigger portion* of the search space
 - So, every once in a while the optimiser "jumps" to some other part of the search space with some probability
- As a *consequence*, it gets to *explore parts* of the search space it would *not have explored otherwise*

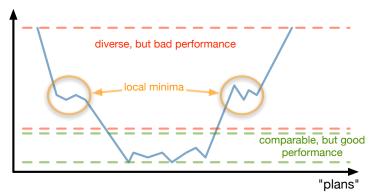
The "well"

cost



The "well" and local minima

cost

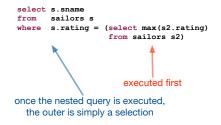


Final step — the entire plan

- The optimiser has produced plans for each query block
- The *question* is now one of *combining* the *sub-plans* to *formulate* the *entire query plan*
- The *strategy* used *depends* on *whether* the *outer* and *nested* queries are *correlated or not*
 - If they are, then in all probability the two sub-plans will be combined through a join

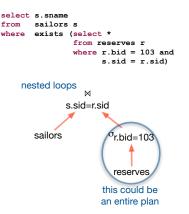
Uncorrelated queries

- Usually, they can be executed in isolation
- The nested query feeds the outer query with results



Correlated queries

- Sometimes, it is *not possible* to *execute* the *nested query just once*
- In those cases the optimiser reverts to a nested loops approach
 - The nested query is executed once for every tuple of the outer query



In practice

- *Before* breaking up the *query into blocks*, most systems *try* to *rewrite* the *query* in some *other way* (*de-correlation*)
 - The idea is that there will probably be a join, so it will be better if the query is optimised in its entirety
- If *de-correlation* is *not possible*, then it is *nested loops all the way*
 - Usually, compute the nested query, store it in a temporary relation and do nested loops with the outer

What do we have and what do we need?

• We have

- A way to decompose a query
- A way to identify equivalent, alternative representations of it (i.e., a search space)
- A statistical framework to estimate cardinalities
- A cost model to estimate the cost of an alternative
- Ways of exploring the search space
- We need
 - Nothing!

Summary

- The *query optimiser* is the *heart* of the *query engine*
 - If it does not do a good job, the engine is doomed to sub-optimal performance
- Two key, closely related decisions
 - Order in which operations are performed
 - Algorithms that perform the operations
- The *paradigm* used is *cost-based optimisation*
 - Three steps: generate alternative plans, estimate the cost of each plan, pick the cheapest
- The *cost model* used is the *cardinality-based* cost model
 - Because cardinality is a good I/O metric
 - ► As a *consequence*, we need *good ways of doing* two things
 - * Estimating the cost of an algorithm
 - ★ Estimating the output cardinality of operations

Summary (cont.)

- Cardinality estimation is 50% of the problem
 - ► Two approaches: uniform distribution assumption, or histograms
 - The uniform distribution assumption essentially does not "care" about the values themselves, they all have an equal probability of appearing
 - Histograms are a better and more elegant distribution approximation technique
 - * Equi-width and equi-depth histograms are the two dominant classes

Summary (cont.)

- The remaining 50% is search space exploration
 - ► Largely *based* on the *equivalence rules* of *relational algebra*
 - Dynamic programming is the dominant approach
 - ★ Find the *cheapest way* to *access single relations*
 - * Find the *cheapest way* to *join two relations*
 - * For each pair, find the cheapest way to join in a third relation
 - ★ Keep going . . .

Summary (cont.)

- Other approaches include rule-based optimisation, randomised exploration, ...
- All approaches aim at one thing
 - Picking a good evaluation plan
 - ► It *might not be* the *cheapest overall*, but it *usually* is of *comparable cost*
- Query optimisation is still an open issue
 - ► We have *good ways* of *solving sub-problems*, but the *entire problem* remains *largely unsolved*

Overview

- So far, we have *focussed* on *query processing*
 - ► In other words, *reading* and *manipulating* data
- A database system, however, not only reads, but also stores data
 - At the same time as others are querying it
- We *need* a way to *ensure concurrent access* to the data
 - Without compromising system performance

Overview

Overview (cont.)

- The basic concept is transaction processing
- Every transaction needs to satisfy four basic properties
 - Atomicity, consistency, isolation, durability
- *How* does the system *guarantee* these *properties*?
 - Remember, without compromising performance
 - Solution: by interleaving transactions

Overview (cont.)

- How can we decide if, after we have interleaved transactions, the result is correct?
 - Interleaving transactions actually causes certain anomalies
 - Solution: the system uses locks to ensure correctness
- How are locks used?
 - Lock granularity, degrees of consistency and two-phase locking
- What *impact* do *locks* have on *performance*?

Overview (cont.)

- Locking poses significant overhead
 - Luckily, however, this overhead can be "tuned" by the user
 - Transaction isolation levels
- But what if the worse comes to worst?
 - System crashes
 - Transactional semantics and recovery
 - Write-ahead logging and the ARIES algorithms

Transactions

- A DBMS spends a lot of time waiting on I/O
 - It is important to keep the CPU busy while waiting
 - In other words, execute other operations concurrently
- *Fact*: the *DBMS* does *not "care" what* the *user does* with the *data* that is *being read* or *written*
 - ► All *it cares about* is that *data* is *being read* or *written*
- A *transaction* is the *DBMS's abstract view* of *user programs*: a *sequence* of *reads* and *writes*

Concurrent execution

- The *transaction user abstraction*: when a *user submits* a *transaction* it is *as if* the *transaction* is *executing by itself*
 - The DBMS achieves concurrency by interleaving transactions
 - ► If the *transaction begins* with the *DB* in a *consistent state*, it *must leave* the *DB* in a *consistent state* after it *finishes*
- The *semantics* of the *transactions* are *unknown* to the *system*
 - Whether the transaction updates a bank account or it fires a rocket missile, the DBMS will never know!

ACID properties

- Atomicity: all the actions in a transaction are executed as a single atomic operation; either they are all carried out or none are
- Consistency: if a transaction begins with the DB in a consistent state, it must finish with the DB in a consistent state
- *Isolation*: a transaction should *execute as if* it is the *only one executing*; it is *protected* (*isolated*) from the *effects* of *concurrently running transactions*
- *Durability*: if a *transaction* has been *successfully completed*, its *effects* should be *permanent*

Example

- Consider two transactions
 - First transaction transfers funds, second transaction pays 6% interest
- If they are submitted at the same time, there is no guarantee as to which is executed first
 - But the end effect should be equivalent to the transactions running serially



Begin A = 1.06*A

End

B = 1.06*B

Example (cont.)

Acceptable schedule

| T1 | A = A+100 | | B = B-100 | |
|----|-----------|------------|-----------|------------|
| T2 | | A = 1.06*A | | B = 1.06*B |

Problematic schedule

| T1 | A = A+100 | | | B = B-100 |
|----|-----------|------------|------------|-----------|
| T2 | | A = 1.06*A | B = 1.06*B | |

DBMS's view

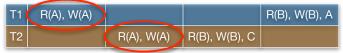
| T1 | R(A), W(A) | | | R(B), W(B) |
|----|------------|------------|------------|------------|
| T2 | | R(A), W(A) | R(B), W(B) | |

Scheduling

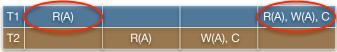
- A schedule is a sequence of reads and writes for some transaction workload incorporating all actions of the workload's transactions
 - ► Serial schedule: the actions of different transactions are not interleaved
 - Equivalent schedules: for any database state, the effect of executing the first schedule is identical to the effect of executing the second schedule
 - Serialisable schedule: a schedule that is equivalent to a serial schedule

Conflicts

Reading uncommitted data (WR conflicts, or "dirty reads")



Unrepeatable reads (RW conflicts)



Overwriting uncommitted data (WW conflicts, or "lost updates")



The solution: locks

- *Before* a *transaction "touches"* a *DB object* it has to *obtain a lock* for it
 - S (Shared) lock for reading
 - X (eXclusive) lock for writing
- Strict two-phase locking (Strict 2PL)
 - Each transaction must obtain an S lock for everything it reads before it starts reading it and an X lock for everything it writes before it starts writing
 - All locks held by a transaction are released only when the transaction commits
 - Once a transaction obtains an X lock for a DB object no other transaction can obtain an X or an S lock for that object
- Strict 2PL produces only serialisable schedules

What can go wrong?

- If a transaction T_i is aborted, then all its actions have to be undone; not only that, but if T_j reads an object written by T_i, T_j needs to be aborted as well (cascading aborts)
- Most systems *avoid cascading aborts* with the following rule:
 - If T_i writes an object T_j can read this object only after T_i commits
- In order to know what needs to be undone, the system keeps a log, recording all writes
- The log is also helpful when recovering from system crashes

The log

- The following actions are recorded in the log
 - Whenever a transaction writes an object
 - * The *log record must* be *on disk before* the *data record* reaches the disk
 - Whenever a transaction commits/aborts
- Log records are chained by transaction ID (why?)
- All *log-related activities* (in fact, all *concurrency control related activities*) are *handled by the DBMS*
 - The user does not know anything

Crash recovery

- Three phases to recovery (ARIES)
 - Analysis: scan log forward, identifying committed and aborted/unfinished transactions
 - ► *Redo*: all *committed transactions* are *made durable*
 - Undo: the actions of all aborted and/or unfinished transactions are undone

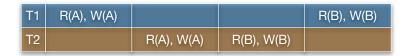
Concurrency control

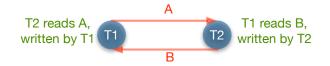
- Serial schedule: the actions of different transactions are not interleaved
- Equivalent schedules: for any database state, the effect of executing the first schedule is identical to the effect of executing the second schedule
- Serialisable schedule: a schedule that is equivalent to a serial schedule
- Two *schedules* are *conflict equivalent* if:
 - ► They *involve* the *same actions* of the *same transactions*
 - Every pair of conflicting actions is ordered the same way
- Schedule S is conflict serialisable if S is conflict equivalent to some serial schedule

Dependency graphs

- Given a schedule S
 - One node per transaction
 - An edge from T_i to T_j , if T_j reads or writes an object written by T_i
- Theorem: a schedule S is conflict serialisable if and only if its dependency graph is acyclic

Example: not conflict serialisable schedule





Review: Strict 2PL

• Strict two-phase locking (Strict 2PL)

- Each transaction must obtain an S (Shared) lock for everything it reads before it starts reading it and an X (eXclusive) lock for everything it writes before it starts writing
- All locks held by a transaction are released only when the transaction commits
- ► Once a transaction obtains an X lock for a DB object no other transaction can obtain an X or an S lock for that object
- Strict 2PL produces only serialisable schedules
 - ► In other words: *schedules* with *acyclic dependency graphs*

Simple 2PL

- Two-phase locking (2PL)
 - Each transaction must obtain an S (Shared) lock for everything it reads before it starts reading it and an X (eXclusive) lock for everything it writes before it starts writing
 - ► A transaction cannot request additional locks once it releases any locks
 - Once a transaction obtains an X lock for a DB object no other transaction can obtain an X or an S lock for that object

Lock management

- Lock and unlock requests are handled by the lock manager that maintains a lock table
- Lock table entry:
 - Number of transactions currently holding a lock
 - Type of lock held (shared or exclusive)
 - Pointer to queue of lock requests
- Locking and unlocking have to be atomic operations
- Lock upgrade: transaction that holds a shared lock can be upgraded to hold an exclusive lock

Deadlocks

- As always, where there are locks, there are deadlocks
- *Deadlocks*: *cycle* of *transactions waiting for locks* to be *released* by *each other*
- Two ways of dealing with deadlocks
 - Deadlock prevention
 - Deadlock detection

Deadlock prevention

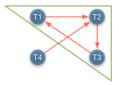
- The *solution* involves *timestamps*; a *timestamp* is the *transaction's priority*
- If *T_i* wants a lock that *T_i* holds, there are two possible policies
 - Wait-Die: if T_i has higher priority, T_i waits for T_j ; otherwise T_i aborts
 - Wound-Wait: if T_i has higher priority, T_j aborts; otherwise T_i waits
- If a transaction re-starts, it has its original timestamp

Deadlock detection

• Create a waits-for graph

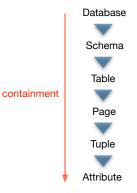
- Nodes are transactions
- There is an edge from *T_i* to *T_j* if *T_i* is waiting for *T_j* to release a lock
- *Periodically check* for *cycles* in the *waits-for graph*

| T1 | S(A) | R(A) | | | S(B) | | | | | |
|----|------|------|------|------|------|------|------|------|------|------|
| T2 | | | X(B) | W(B) | | | | X(C) | | |
| T3 | | | | | | S(C) | R(C) | | | X(A) |
| T4 | | | | | | | | | X(B) | |



Multiple granularity locks

- What should we lock? Tuples, pages, tables, ...
- But there is an *implicit* containment
- *Idea*: *lock* DB objects *hierarchically*



Hierarchical locks and new locking modes

- Allow transactions to lock at each level of the hierarchy
- Introduce "intention" locks: IS and IX
 - Before locking an item, a transaction must introduce intention locks on all the item's ancestors in the hierarchy
 - Release locks in reverse order
- One extra lock: SIX "share, with intention to write"

Compatibility matrix

| | | NL | IS | IX | SIX | S | Х |
|-------------|-----|----|----|----|-----|---|---|
| ock | NL | Y | Y | Y | Y | Y | Y |
| | IS | Y | Y | Y | Y | Y | N |
| wanted lock | IX | Y | Y | Y | Ν | Ν | N |
| war | SIX | Y | Y | Ν | Ν | Ν | N |
| | S | Y | Y | Ν | Ν | Y | N |
| | Х | Y | Ν | Ν | Ν | Ν | Ν |

held lock

In more detail

- Each transaction starts from the root of the hierarchy
- To obtain S or IS lock on a node, must hold IS or IX on parent node
 - What if a transaction holds SIX on parent? S on parent?
- To obtain X or IX or SIX on a node, must hold IX or SIX on parent node
- Must release locks in bottom-up order

A few examples

- T1 scans R, and updates a few tuples
 - ► T1 gets an SIX lock on R, then repeatedly gets an S lock on tuples of R, and occasionally upgrades to X on the tuples
- T2 uses an index to read only part of R
 - ► T2 gets an IS lock on R, and repeatedly gets an S lock on tuples of R
- T3 reads all of R
 - T3 gets an S lock on the entire relation
 - ► Or, it gets an IS lock on R, escalating to S lock on every tuple

Here's the catch (the phantom problem)

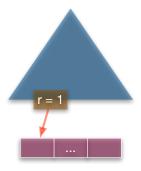
- If we *relax* the *assumption* that the *DB* is a *fixed collection* of objects, *even Strict 2PL* will *not assure serialisability*!
 - ► T1 locks all pages containing sailor records with rating = 1, and finds oldest sailor (say, age = 71)
 - Next, *T2 inserts* a *new sailor*: rating = 1, age = 96
 - ► T2 also deletes oldest sailor with rating = 2 (and, say, age=80), and commits
 - T1 now locks all pages containing sailor records with rating = 2, and finds oldest (say, age=63)
- No lock conflicts, but also no consistent DB state where T1 is "correct"!

The problem

- *T1 implicitly assumes* that it has *locked the* set of *all sailor* records with *rating* = 1
 - The assumption only holds if no sailor records are added while T1 is executing!
 - ▶ We need some mechanism to enforce this assumption
 - ★ Index locking
 - ★ Predicate locking
- The example shows that conflict serialisability guarantees serialisability only if the set of objects is fixed!

Index locking

- If there is an index on the rating field, T1 should lock the index page containing the data entries with rating = 1
 - If there are no records with rating = 1, T1 must lock the index page where such a data entry would be, if it existed!
- If there is no suitable index, T1 must lock all pages, and lock the file/table to prevent new pages from being added, to ensure that no new records with rating = 1 are added



Predicate locking

- Grant lock on all records that satisfy some logical predicate, e.g., salary > 2 · salary
 - Index locking is a special case of predicate locking for which an index supports efficient implementation of the predicate lock
 - What is the predicate in the sailor example?
- In general, predicate locking imposes a lot of locking overhead

B+tree locking

- How can we efficiently lock a particular node?
 - This is entirely different than multiple granularity locking (why?)
- One *solution*: *ignore* the *tree structure*, just *lock pages* while *traversing* the tree, following *2PL*
 - Terrible performance
 - Root node (and many higher level nodes) become bottlenecks because every tree access begins at the root

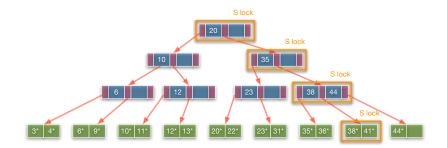
Key observations

- Higher levels of the tree only direct searches to leaf pages
- For *insertions*, a *node* on a *path* from the *root* to a modified *leaf* must be *locked* (in *X mode*, of course), *only if* a *split* can *propagate up* to it *from* the *modified leaf* (similar point holds for deletions)
- We can *exploit* these *observations* to design *efficient locking protocols* that *guarantee serialisability* even though they *violate 2PL*

The basic algorithm

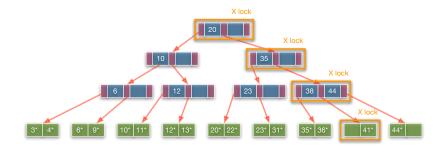
- Search: start at root and descend; repeatedly, S lock child then unlock parent
- *Insert/Delete*: *start* at *root* and *descend*, obtaining X *locks as needed*; once *child* is *locked*, *check* if it is *safe*:
 - Safe node: a node such that changes will not propagate up beyond this node
 - * Insertion: node is not full
 - ★ Deletion: node is not half-empty
 - ► If *child* is *safe*, *release* all *locks* on *ancestors*

Example: search 38*



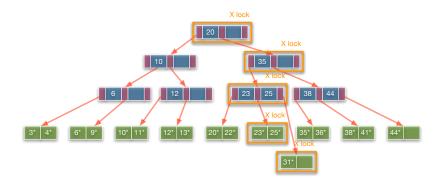
Obtain and release S-locks level-by-level

Example: delete 38*



Obtain X-locks while descending; release them top-down once the node is designated safe

Example: insert 25*

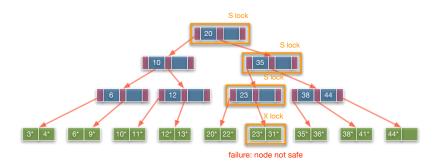


Obtain X-locks while descending; leaf-node is not safe so create a new one and lock it in X-mode; first release locks on leaves and then the rest top-down

Optimistic B+tree locking

- Search: as before
- *Insert/delete*: set *locks as* if for *search*, get to the leaf, and *set X lock* on the leaf
 - If the leaf is not safe, release all locks, and restart transaction, using previous insert/delete protocol
- "Gambles" that only leaf node will be modified; if not, S locks set on the first pass to leaf are wasteful
 - In practice, better than previous algorithm

Example: insert 25*

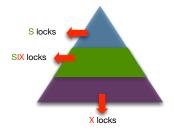


Obtain S-locks while descending, and X-lock at leaf; the leaf is not safe, so abort, release all locks and restart using the previous algorithm

Even better algorithm

- Search: as before
- Insert/delete: use original insert/delete protocol, but set IX locks instead of X locks at all nodes
 - Once leaf is locked, convert all IX locks to X locks top-down: i.e., starting from the unsafe node nearest to root
 - Top-down reduces chances of deadlock
 - * Remember, this is *not the same* as *multiple granularity locking*!

Hybrid approach



- The *likelihood* that we will need an X lock decreases as we move up the tree
- Set *S locks* at *high levels*, *SIX locks* at *middle levels*, *X locks* at *low levels*

Transaction isolation

| Isolation level | Dirty read | Unrepeatable read | Phantoms |
|---------------------|------------|----------------------|----------|
| Read uncommitted | Maybe | Maybe | Maybe |
| Read committed | No | Maybe | Maybe |
| Repeatable reads | No | No | Maybe |
| Serialisable | No | No | No |

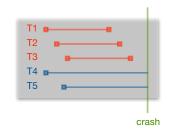
Review: ACID properties

- Atomicity: all the actions in a transaction are executed as a single atomic operation; either they are all carried out or none are
- Consistency: if a transaction begins with the DB in a consistent state, it must *finish* with the *DB* in a *consistent state*
- *Isolation*: a transaction should *execute as if* it is the *only one* executing; it is protected (isolated) from the effects of concurrently running transactions
- Durability: if a transaction has been successfully completed, its effects should be *permanent*

Atomicity and durability are ensured by the recovery algorithms

What can go wrong?

- Atomicity
 - Transactions may abort; their effects need to be undone
- Durability
 - What if the system stops running?



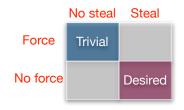


Problem statement

- Updates are happening in place
 - There is a *buffer pool*
 - ★ Data pages are read from disk
 - * Data pages are modified in memory
 - ★ Overwritten on, or deleted from disk
- We need a *simple scheme* to *guarantee atomicity* and *durability*

More on the buffer pool

- Two issues: force and steal
- *Force*: when a *data page* is *modified* it is *written* straight to *disk*
 - Poor response time
 - But durable
- *Steal*: effects of *uncommitted transactions* reach the *disk*
 - Higher throughput
 - But not atomic



The problems

- *Steal*'s problems are all about *atomicity*
 - What if a transaction modifying a page aborts?
 - If we steal a page, we need to remember its old value so it can be restored (UNDO)
- *No force*'s problems are all about *durability*
 - What if a system crashes before a modified page is written to disk?
 - We need to record enough information to make the changes permanent (REDO)

Recovery

The solution: logging

- *Record REDO* and *UNDO* information in a record of a *separate* structure: the log
 - Sequential writes for every update
 - Minimal information written (more efficient!)
 - Keep it on a separate disk!
- Log: a list of REDO and UNDO actions
 - Each log record contains at least:
 - * Transaction id, modified page, old data, new data

Recovery

Write-ahead logging

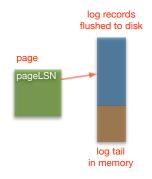
- The log adheres to the write-ahead protocol (WAL)
 - Must force the log record for an update before the corresponding data page gets to disk
 - Must force all log records for a transaction before it commits
- #1 guarantees *atomicity*
- #2 guarantees durability

Normal execution

- Series of reads and writes
- Followed by a commit (success) or abort (failure)
- Steal, No-force management
- Adherence to the WAL protocol
- Checkpoints: periodically, the system creates a checkpoint to minimise the time taken to recover
 - Assume the DB is consistent after a checkpoint

WAL and the log

- Each *log record* has a unique *log* sequence number (LSN)
 - LSNs are always increasing
- Each *data page* contains a *pageLSN*
 - The LSN of the most recent log record for an update to that page
- The *system* keeps track of flushedLSN
 - ► The max LSN flushed so far
- WAL: before a page is written, pageLSN ≤ flushedLSN



Recovery

Log records

• Possible *log records types*

- Update
- ► Commit
- Abort
- End (signifies commit or abort!)
- Compensation Log Records (CLR)
 - ★ Logging UNDO actions!
 - ★ But we will not talk about them in more detail



Other log-related state

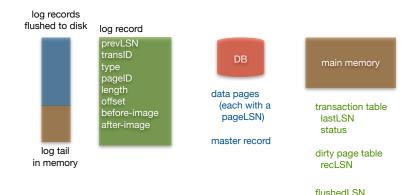
- Transaction table: one entry per active transaction
 - Contains transaction id, status (running/commited/aborted) and lastLSN — log sequence number of the last log record for that transaction
- Dirty page table: one entry per dirty page in buffer pool
 - Contains recLSN the LSN of the log record which first caused the page to be dirty

Recovery

Checkpoint records

- **begin_checkpoint** record: indicates when checkpoint began
- end_checkpoint record: contains *current transaction table* and *dirty* page table
- This is a *"fuzzy checkpoint"*
 - Other transactions continue to run; so these tables accurate only as of the time of the begin_checkpoint record
 - No attempt to force dirty pages to disk; effectiveness of checkpoint limited by oldest unwritten change to a dirty page
 - So it's a good idea to periodically flush dirty pages to disk
- Store LSN of checkpoint record in a safe place (master record)

What's stored where



Simple transaction abort

- For now, consider an *explicit abort* of a transaction
 - No crash involved
- We want to "*play back*" the log in *reverse order*, UNDO ing updates
 - Get last SN of transaction from transaction table
 - Follow chain of log records backward via the prevLSN field
 - Before starting UNDO, write an Abort log record
 - ★ For recovering from crash during UNDO!

Abort (cont.)

- To perform UNDO, must have a lock on data
 - No problem
- Before restoring old value of a page, write a CLR
 - Continue logging while you UNDO!
 - CLR has one extra field: undonextLSN
 - ★ Points to the next LSN to undo (i.e., the prevLSN of the record we're currently undoing)
 - CLRs are never undone (but they might be redone when repeating history: guarantees atomicity)
- At the end of UNDO, write an "end" log record

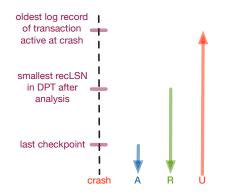
Recovery

Transaction commit

- Write *commit record* to log
- All log records up to the transaction's lastLSN are flushed
 - ► Guarantees that flushedLSN ≥ lastLSN
 - Note that log flushes are sequential, synchronous writes to disk
 - Many log records per log page
- Commit() returns
- Write end record to log

Recovery

Recovery: big picture



- *Start* from a *checkpoint* (found via *master record*)
- Three phases
 - Analysis: figure out which transactions committed since the checkpoint, and which failed
 - REDO all actions
 - ★ Repeat history
 - UNDO effects of failed transactions

Additional issues

- What happens if the system crashes during the analysis phase? During *REDO phase*?
- How can the *amount of work* during *REDO* be *limited*?
 - Flush asynchronously in the background
- How can the amount of work during UNDO be limited?
 - Avoid long-running transactions

Summary

- Concurrency control and recovery are key concepts of a DBMS
- *Both* are *ensured* by the *system itself*; the user does not (and should not!) know of their existence
- The key abstraction is the transaction
 - The processing unit of the system
 - Four key properties
 - * Atomicity, consistency, isolation, durability

Summary (cont.)

- A transaction is viewed by the system as a series of reads and writes
- To *improve throughput*, the *system interleaves* the *actions* of the *transactions* (*i.e.*, a schedule)
 - ► At all times, *ensuring serialisability* of the *produced schedules*
- Locks are the mechanism that ensures serialisability
 - *Before reading*, obtain a *Shared lock*
 - Before writing, obtain an eXclusive lock

Summary (cont.)

- Multiple granularity of locks
 - Leads to an escalation of locks, as we are descending the hierarchy
- Special protocols for indexes and predicates
- Transactions help after recovering from a crash
 - ▶ As the processing unit, we know what needs to be repeated or deleted

Summary (cont.)

- Steal, no-force buffer pool management
 - Higher response time (steal)
 - Higher throughput (no-force)
- Need to use it, without satisfying correctness
- Use a log to record all actions
 - Employ the Write-Ahead Logging protocol

Summary (cont.)

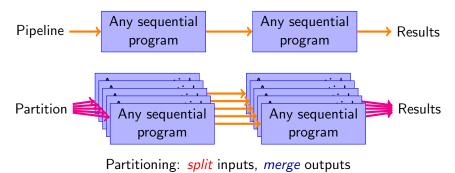
- Use *checkpoints* to *periodically record consistent states* and *limit* the amount of the *log* that needs to be *scanned during recovery*
- *Recovery* in *three phases*
 - ► Analysis: from checkpoint, figure out REDO and UNDO extents
 - REDO: repeat entire history
 - UNDO: delete effects of failed transactions
- Repeating history simplifies the logic

Why parallelism?

- The very *definition* of *parallelism*: *divide* a big *problem* into many *smaller* ones to be *solved in parallel*
- Consider we have a *terabyte* of data to *scan*
 - ▶ With *one pipe* of 10*MB/s*, we need 1.2 *days*
 - By partitioning the data in disjoint subsets and having 1,000 parallel pipes of the same bandwidth, we need 90s

Parallelism and DBMSs

- Parallelism is natural to DBMS processing
 - Pipeline parallelism: many machines each doing one step in a multi-step process
 - Partition parallelism: many machines doing the same thing to different pieces of data.
 - Both are natural in a DBMS



The parallelism success story

• DBMSs are the most (only?) successful application of parallelism

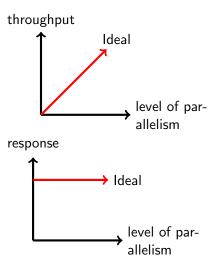
- ► Teradata, Tandem vs. Thinking Machines, KSR, ...
- Every major DBMS vendor has some parallel server
- Workstation manufacturers now depend on parallel DB server sales
- *Reasons* for success
 - Bulk-processing (partition parallelism)
 - Natural pipelining
 - Inexpensive hardware can do the trick
 - Users/app-programmers do not need to think in parallel

Terminology

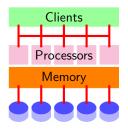
Speed-up

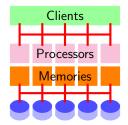
More resources means proportionally less time for given amount of data (throughput)

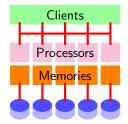




Architecture: what to share?







Shared memory

- Easy to program
- Expensive to build
- Difficult to scale up

Shared disk

- *Middle* of the road
- *Distributed* file system
- *Cluster* computing

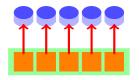
Shared nothing

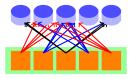
- Hard to program
- Cheap to build
- Easy and ideal to speed/scale up

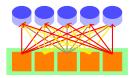
Different types of parallelism

- Intra-operator parallelism
 - All machines working to compute a single operation (scan, sort, join)
- Inter-operator parallelism
 - Each operator may run concurrently on a different site (exploits pipelining)
- Inter-query parallelism
 - Different queries run on different sites
- We shall focus on intra-operator parallelism

Automatic data partitioning







Range

- Good for equi-joins
- Range-queries

• Good for aggregation

Hash

- Good for equi-joins
- No range-queries
- Problematic with skew

Round-robin

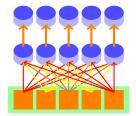
- Indifferent for equi-joins
- Range-queries complicated
- Load-balanced

Parallel scans

- Scan in parallel, and merge
- Selections may not require all sites for range or hash partitioning
- Indexes can be built at each partition
- Question: how do indexes differ in the different schemes?
 - Think about both lookups and inserts!
 - What about key indexes?

Parallel sorting

- Key idea: sorting *phases* are intrinsically *parallelisable*
 - Scan in parallel, range-partition as you go
 - As tuples come in, begin "local" sorting using standard algorithm
 - Resulting data is sorted, and range-partitioned
- Problem: *skew*
 - Solution: sample the data to determine partition points



Parallel aggregation

- For each aggregate function, need a decomposition
 - $count(S) = \sum_{i} count(s(i))$, ditto for sum()
 - $avg(S) = (\sum_{i} sum(s(i))) / \sum_{i} count(s(i))$
 - and so on . . .
- For groups
 - Sub-aggregate groups close to the source
 - Pass each sub-aggregate to its group's site
 - ★ Chosen via a *hash function*

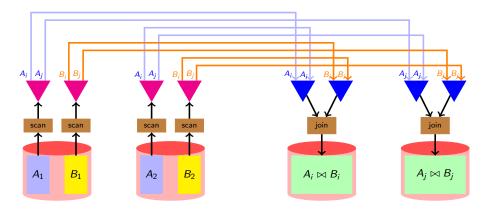
Parallel joins

- Nested loops
 - Each outer tuple must be compared with each inner tuple that might join
 - ► Easy for range partitioning on join columns, hard otherwise
- Sort-merge (or plain merge-) join
 - Sorting gives range-partitioning
 - Merging partitioned tables is local

Parallel hash join

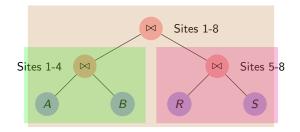
- During the *first phase*, *partitions* are *distributed* to *different sites*
 - A good hash function automatically distributes work evenly
- Second phase is local at each site
 - Almost always the winner for equi-join
- Good use of split/merge makes it easier to build parallel versions of sequential join code

Dataflow network for parallel join



Complex parallel query plans

- Complex queries: inter-operator parallelism
 - Pipelining between operators
 - * Note that sorting and phase one of hash-join block the pipeline (yet again!)
 - Bushy execution trees



Observations

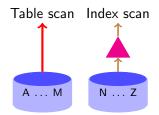
- It is relatively easy to build a fast parallel query executor
- It is hard to write a robust and world-class parallel query optimizer
 - There are many tricks
 - One quickly hits the *complexity barrier*
 - Still open research

Parallel query optimization

- Common approach: two phases
 - Pick best sequential plan (System R algorithm)
 - Pick degree of parallelism based on current system parameters
- Allocate operators to processors
 - ► Take *query tree*, *decorate* as in previous example

What can go wrong?

- Best sequential plan \neq best parallel plan
- Trivial counter-example
 - Table partitioned with local secondary index at two nodes
 - ► *Range query*: all of node 1 and 1% of node 2
 - ★ e.g., select * from telephone_book where name < "NoGood"
 - Node 1 should do a scan of its partition
 - Node 2 should use secondary index



Parallel databases summary

- Parallelism natural to query processing
 - Both pipeline and partition parallelism
- Shared-nothing vs. Shared-memory
 - Shared-disk too, but less standard
 - Shared-mem easy, costly; does not scaleup
 - Shared-nothing cheap, scales well, harder to implement
- Intra-operator, inter-operator, and inter-query parallelism all possible.

Parallel database summary (cont.)

- Data layout choices important
- Most database operations can be done using partition-parallelism
 - Sort
 - Sort-merge join, hash-join
- Complex plans
 - ► Allow for *pipeline-parallelism*, but sorts, hashes *block* the *pipeline*
 - Partition-parallelism achieved through bushy trees

Parallel database summary (cont.)

• Hardest part: optimization

- ► *Two-phase* optimization *simplest*, but can be *ineffective*
- More complex schemes still at the research stage
- We have not discussed transactions, logging
 - Easy in shared-memory/shared-disk architecture
 - Takes some care in shared-nothing
 - Some ideas from distributed transactions are handy