#### Advanced Databases

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## Outline

## 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

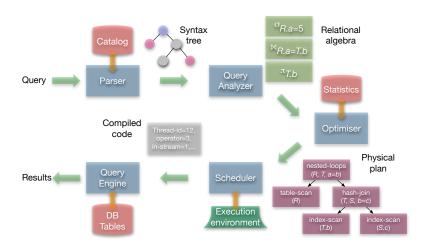
- Introduction
- Relational databases overview
  - ► Data model, evaluation model
- Storage
  - ► Indexes, multidimensional data
- Query evaluation
  - ▶ Join evaluation algorithms, execution models
- Query optimisation
  - ► Cost models, search space exploration, randomised optimisation
- Concurrency 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



#### Outline

## Three basic building blocks

Attribute

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

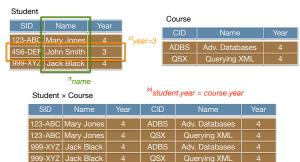


SID	Name	 Year
123-ABC	Mary Jones	 4

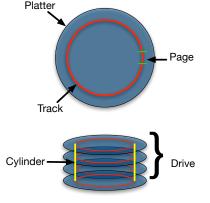
SID	Name	 Year
123-ABC	Mary Jones	 4
456-DEF	John Smith	 3
999-XYZ	Jack Black	4

#### Data manipulation

- Operations to *isolate* a *subset* of a *single relation*: Selection  $(\sigma)$ , Projection  $(\pi)$
- 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 <u>dimensions</u> of the <u>head</u> are <u>impressive</u><sup>1</sup>. With a <u>width</u> of less than a <u>hundred nanometers</u> and a <u>thickness</u> of about <u>ten</u>, it flies above the platter at a <u>speed</u> of up to <u>15,000 RPM</u>, at a <u>height</u> that is the equivalent of <u>40 atoms</u>. 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

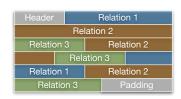
<sup>&</sup>lt;sup>1</sup>Source: Matthieu Lamelot. Tom's Hardware.

# What about flash memory and solid state?

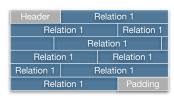
- The *geometry* is different
  - ▶ There are no tracks, or platters, or cylinders or anything of the sort
- But the *issues* are *similar* 
  - Data is still accessed in blocks
  - Blocks are still organised in pages
  - ► Sequential vs. random I/O is still a problem
- Most of the things we say in this 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



Interleaved tuples



Clustered tuples

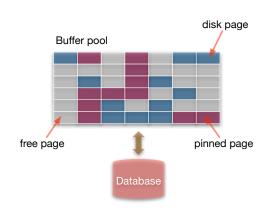
# Advantages of clustering

- 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)

## Outline

# 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<sup>-8</sup> seconds; Comparing two values on disk costs 18.2 · 10<sup>-5</sup> seconds (3 orders of magnitude more expensive.)

## Outline

## 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

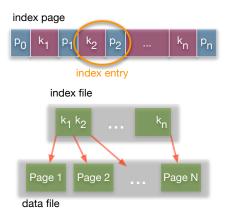
- Tree-structured indexes
  - Much like you would use a binary tree to search, but with a higher key-per-node cardinality
  - Retains order
  - ► Great for *range queries*
  - ▶ Both *one*-dimensional and *multi*-dimensional
- Hash-based indexes
  - ► Fully *randomized* (i.e., no order)
  - ► Great for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

#### Outline

#### Sorted indexes

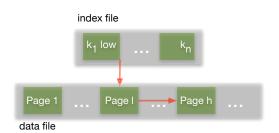
#### • The basic idea:

- 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)



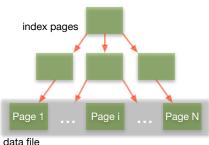
# How does it answer range queries?

- 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!

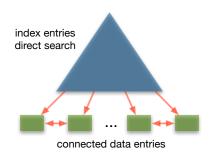


# Potential problem (and the solution)

- 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!

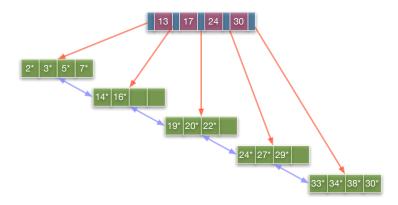


#### B+trees: the most widely used indexes



- Insertion/deletion at log<sub>f</sub> 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

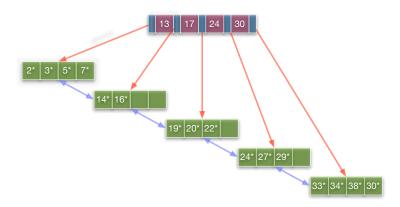


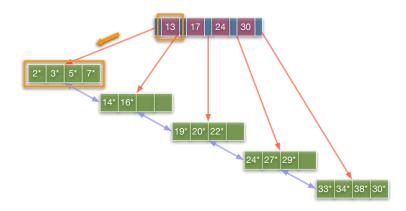
## B+trees in practice

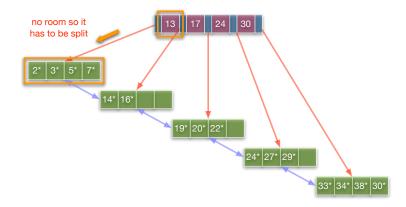
- Typical *order*: 100, typical *fill-factor*: 67%
  - ► Average fan-out: 133
- Typical capacities
  - ► Height 3: 2,532,637
  - ► Height 4: 312,900,700 (!)
- The top levels can often be kept in memory
  - ▶ 1st level: 4,096, or 8,192 bytes (1 page)
  - ▶ 2nd level: 0.5, or 1MB (133 pages)
  - ▶ 3rd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

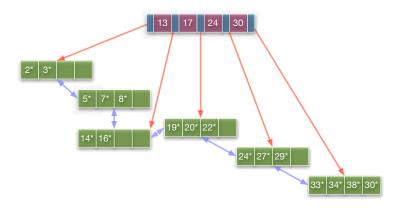
#### B+tree insertion

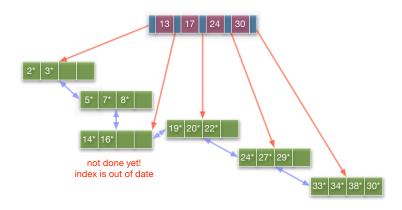
- Find correct leaf L
- Put data entry into L
  - ▶ *If* there is *enough space* in *L*, *done*!
  - ▶ If there is no space, L needs to be split into L and L'
  - Redistribute entries evenly in L and L'
  - ► Insert index entry pointing to L' into the parent of L
- Ascend the tree recursively, splitting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



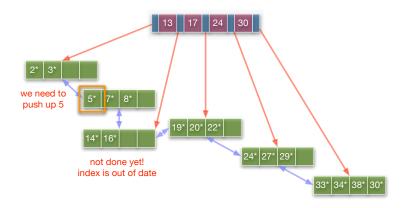




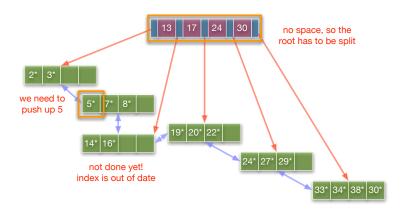




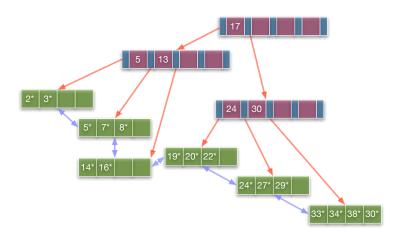
#### B+tree insertion: 8\*



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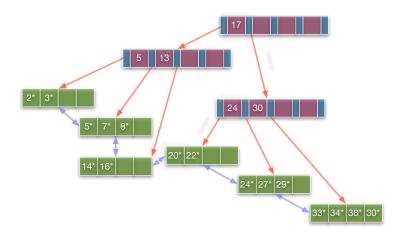
#### Insertion observations

- 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?)

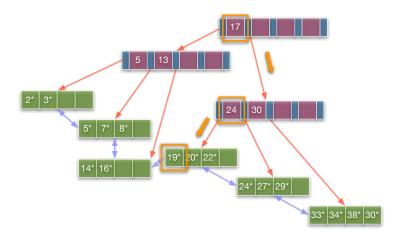
#### B+tree deletion

- 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

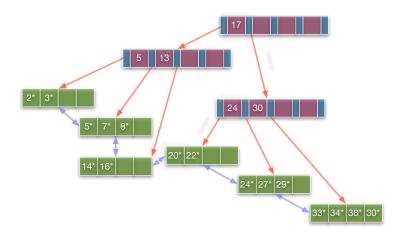
#### B+tree deletion: 19\*

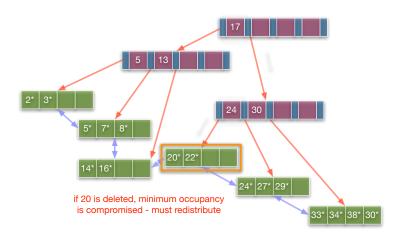


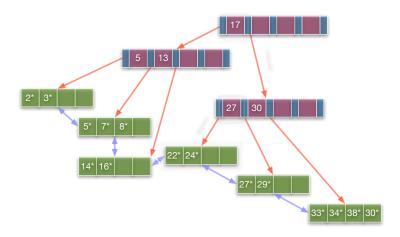
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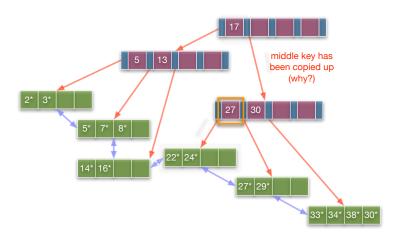


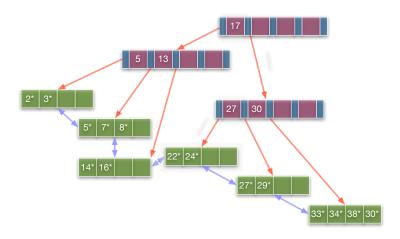
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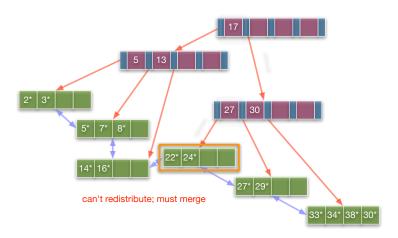


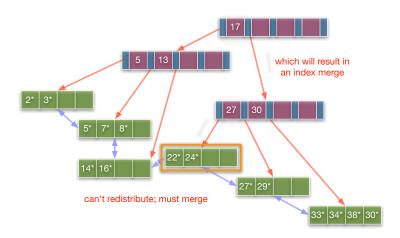


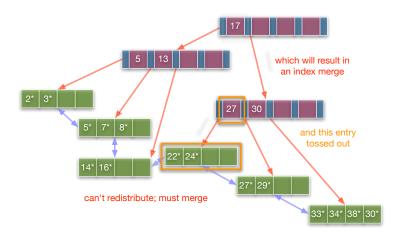




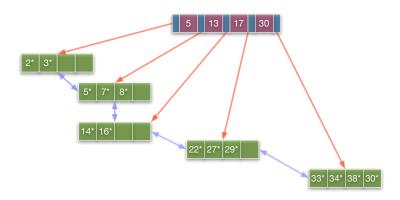








### B+tree after deletion of 24\*

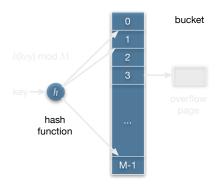


### Summary of B+tree indexes

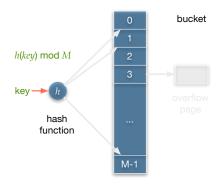
- Ideal for range searches, good for equality searches
- Highly dynamic structure
  - ► Insertions and deletions leave tree height-balanced, log<sub>f</sub> N cost
  - ► For most *typical implementations*, *height* is *rarely greater* than *3 or 4*, occupancy at 67%
  - Which means that the index is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

#### Hash indexes

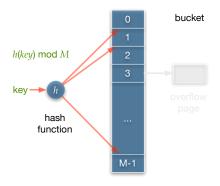
- Hash-based indexes are good for equality selections, not for range selections
  - ▶ In fact, they *cannot support range* selections (why?)
- Static and dynamic techniques exist here as well
  - ► Trade-offs similar to those between ISAM and B+trees



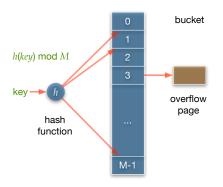
- Number of primary pages fixed
  - Allocated sequentially, never de-allocated
  - Overflow pages if needed
- h(k) mod M = bucket to which data entry with key k belongs (M = number of buckets)



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## Static hashing observations

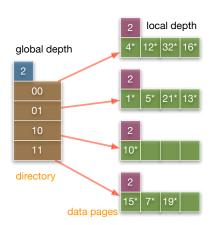
- The buckets contain the actual data!
  - But only the key is hashed
  - No secondary index like in the tree case
- The hash function must uniformly distribute the keys across all buckets
  - Lots of ways to tune the hash function
- Again, long overflow chains of pages will develop, and pretty soon 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

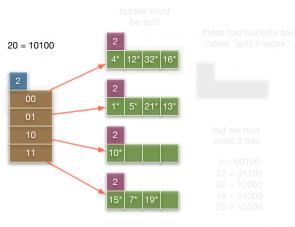
## Extendible hashing example

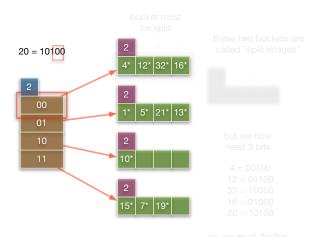
- Directory: array of size 4
- Key k, apply hash function h(k) and translate the result to binary
  - e.g., h(k) = 5 = 101
- Last global depth number of bits identify the bucket



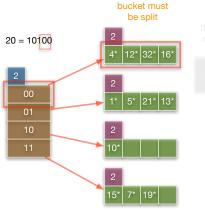
## Global, local depth and doubling

- 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



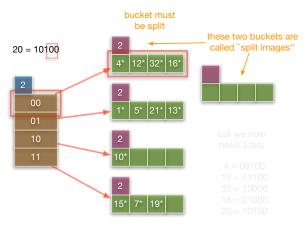


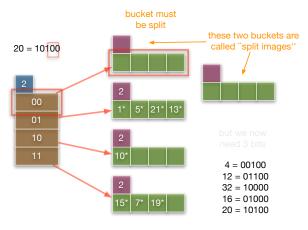
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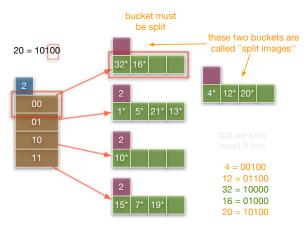


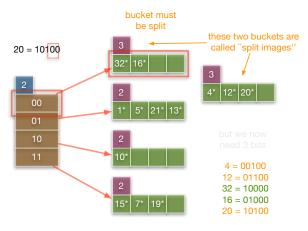
these two buckets are called "split images"

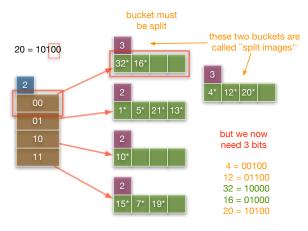
but we now need 3 bits



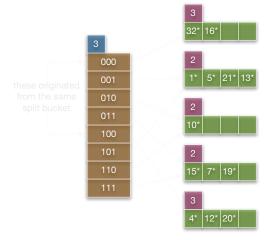




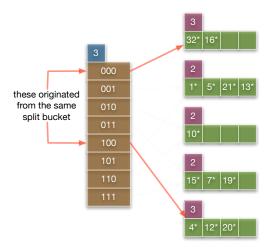




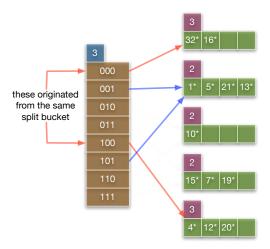
## Doubling the directory



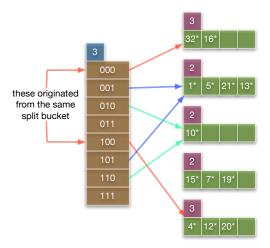
# Doubling the directory



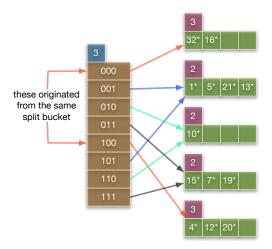
### Doubling the directory



## Doubling 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  $h_0, h_1, h_2, ...$ 
  - $h_i(key) = g(key) \mod (2^i M)$
  - ► *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  $d_i = d_0 + i$ .
  - $\blacktriangleright$   $h_{i+1}$  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<sub>R</sub> initial (for round R) buckets are split
  - ▶ Buckets 0 to N-1 have been split
  - ▶ Buckets N to M<sub>R</sub> 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.

#### Search and insert

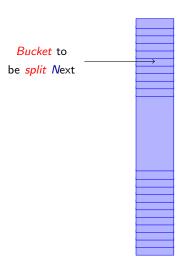
#### Search

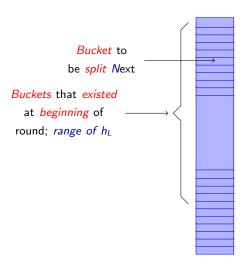
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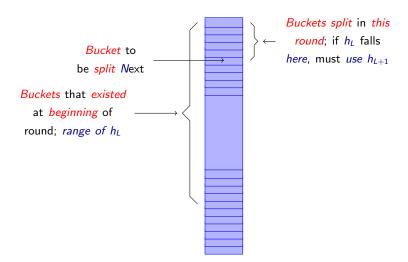
#### Insert

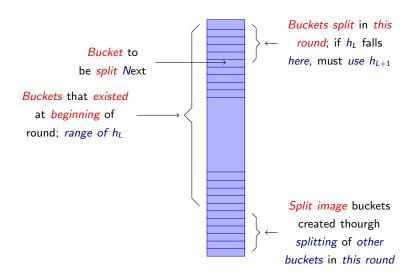
- Find bucket as above, by applying  $h_L$  or  $h_{L+1}$
- If bucket to insert is full
  - Add overflow page and insert entry
  - ► (Maybe) Split bucket N and increment N



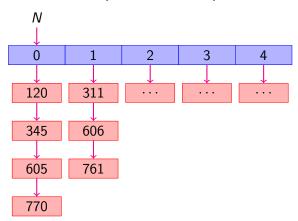








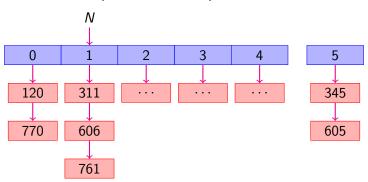
### Splitting a bucket (0 in this case)



#### Hash functions

•  $h_0(K) = K \mod 5$ 

### Splitting a bucket (0 in this case)



#### Hash functions

- $h_0(K) = K \mod 5$
- $h_1(K) = K \mod 10$

# Algorithms in more detail

```
Lookup for key K

bucket := h_L(K);

if bucket < N then bucket = h_{L+1}(K)
```

## Algorithms in more detail

```
Lookup for key K
 bucket := h_I(K);
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```

#### Expansion

```
N := N + 1:
if N = M2^L then
 L := L + 1; N := 0;
```

## Algorithms in more detail

```
Lookup for key K

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### Expansion

```
N := N + 1;
if N = M2^L then
L := L + 1; N := 0;
```

#### Contraction

```
N := N - 1;
if N < 0 then
L := L - 1; N := M2^{L} - 1;
```

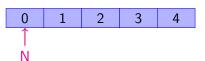
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## Expansion

N := N + 1:

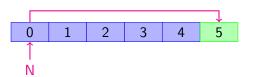
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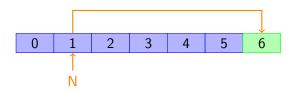


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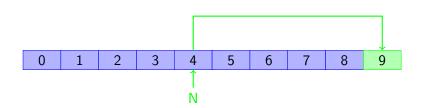
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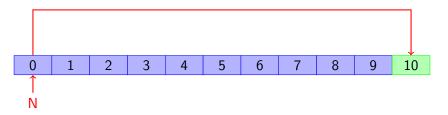
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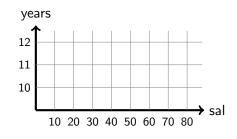
### Linear hashing observations

- 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

#### Outline

### Why more than one dimensions?

- Single-dimensional indexes are not enough
  - ► Consider a *composite search* key *e.g.*, an index on ⟨*sal*, *years*⟩

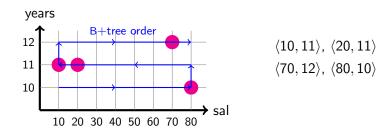


 $\langle 10,11 \rangle$ ,  $\langle 20,11 \rangle$ 

 $\langle 70, 12 \rangle$ ,  $\langle 80, 10 \rangle$ 

### Why more than one dimensions?

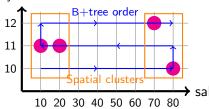
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  - ► We *sort* entries *first* by *sal* and *then* by *years*



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- 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

# years

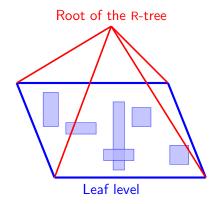


$$\langle 10, 11 \rangle$$
,  $\langle 20, 11 \rangle$ 

$$\langle 70, 12 \rangle$$
,  $\langle 80, 10 \rangle$ 

#### The R-tree

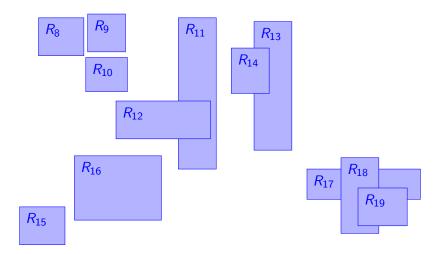
- The R-tree is a tree-structured index that remains balanced on insertions and deletions
- Each key stored in a leaf entry is intuitively a box, or collection of intervals, with one interval per dimension



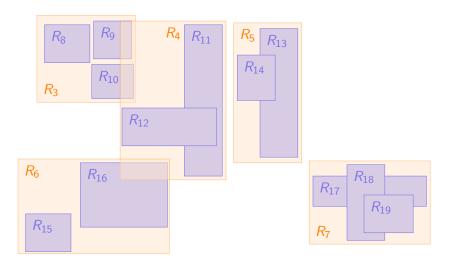
#### R-tree properties

- 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

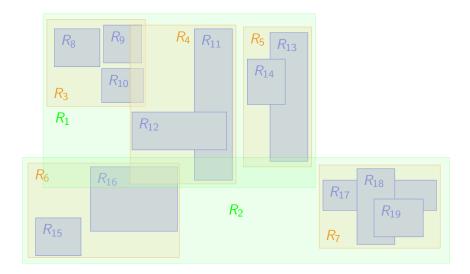
### R-tree example



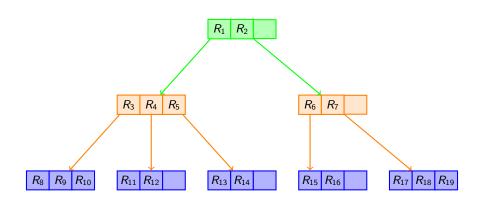
#### R-tree example



### R-tree example



# R-tree example (cont.)



Start at root



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If current node is non-leaf

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

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

```
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```

```
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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* 

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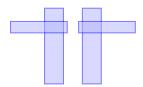
Go to *child* whose *box* needs *least enlargement* to *cover B*; *resolve ties* by going to *smallest area child* 

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.)

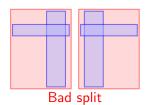
## Splitting a node

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



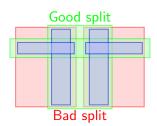
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#### Redistribution

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.

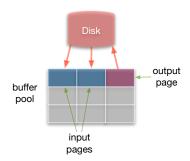
### Outline

#### Overview

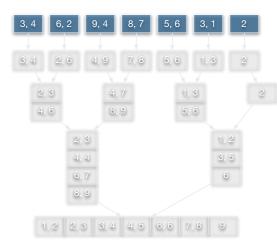
- Sorting is probably the most classic problem in CS
  - ► Simple idea: impose a total order on a set of values
- It is a *classic problem* in *databases* too
  - ► Remember *ISAM*? First step is to sort the file
  - In fact, if you're bulk loading a B+tree, you're better off sorting the file first
- 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

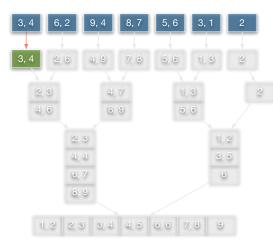
- Requires a *maximum of three* buffer pages and multiple passes over the data
- First pass: read one page, sort it, write it out
- Subsequent passes: read two pages, merge them, write out the result



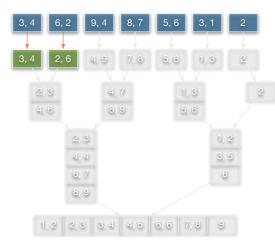
- Each pass will read and write each page in the file
- *N pages*, so the number of passes is  $\lceil \log_2 N \rceil + 1$
- So, the *total I/O cost* is  $2N(\lceil \log_2 N \rceil + 1)$



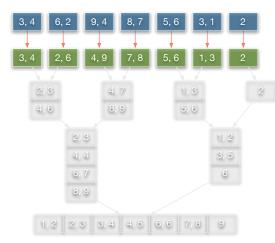
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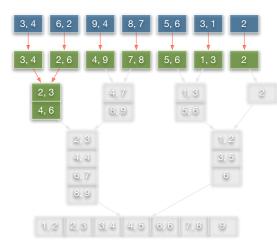
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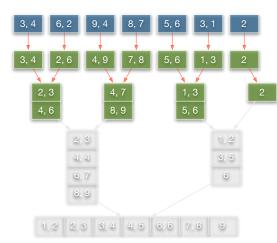
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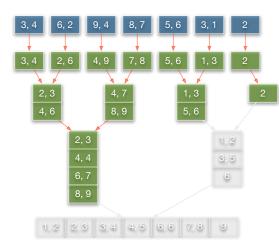
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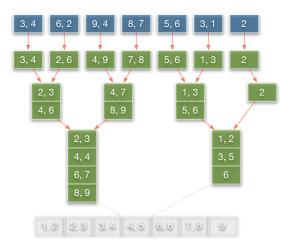
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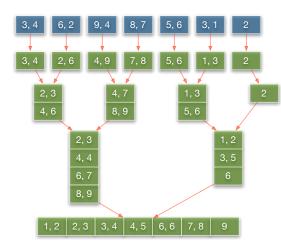
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## But why only three pages?

- 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  $(\lceil \frac{N}{B} \rceil)$
  - ▶ Subsequent passes: merge B 1 runs (why?)

• Number of passes:  $1 + \lceil \log_{B-1} \lceil \frac{N}{B} \rceil \rceil$ 

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    - ► Pass 3: final merge, done!

# A bit of perspective

N	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

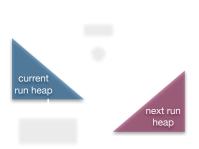
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		257 * 4,096 = 1,052,672					
N	B=3	B=5	B=9	B=17	B=129	B=257	
100	7	4	3	2	1	1	
1,000	10	5	4	3	2	2	
10,000	13	7	5	4	2	2	
100,000	17	9	6	5	3	3	
1,000,000	20	9	7	5	3	3	
10,000,000	23	12	8	6	4	3	
100,000,000	26	14	9	7	4	4	
1,000,000,000	30	15	10	8	5	4	

#### Are we done?

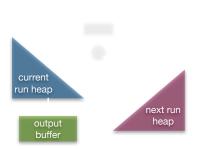
- No! We can actually do much better than this
- Key observation: we are using main memory algorithm (e.g., quicksort) to sort pages in memory
  - ▶ But that *doesn't minimise I/O*, does it?
  - Wouldn't it be nice if we could generate sorted runs longer than memory?
  - ► Solution: heapsort (a.k.a.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

### How does heapsort work?



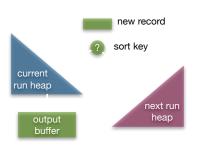
- Keep two heaps in memory, one for each run (the current and the next one)
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- Keep adding to the current run until we are out of buffer space
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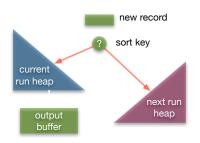
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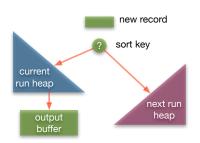
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while (not finished) do {
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```
Initialisation: read B pages into the current heap
while (not finished) do {
  while (r = lowest key from current heap) {
    write r to the current run
    max = r
    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
```

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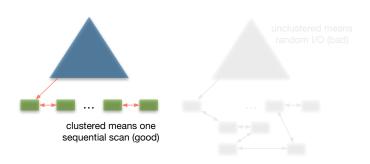
#### Heapsort observations

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

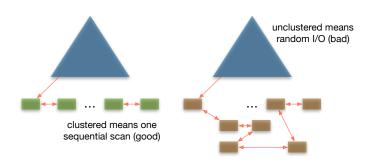
#### Good-old B+trees

- 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



## Clustered vs. unclustered storage



## Summary of sorting

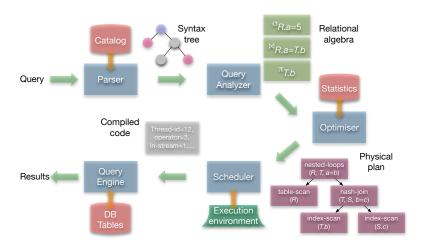
- Databases spend a lot of their time sorting
- In fact, they might dedicate part of their buffer pool for sorting data
  - ▶ Remember pinning buffer pool pages?
- External sorting minimises I/O cost
  - ► First you produce sorted runs, then you merge them
- The choice of internal sort matters as well
  - ► 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

#### Outline

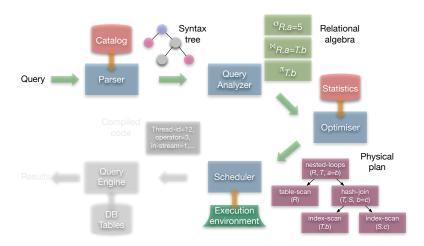
#### Overview

- 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



## Query cycle



#### Outline

#### 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

## Example

#### Algebraic expression

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

#### Example

#### SQL query

#### Algebraic expression

```
\pi_{student.id,student.name}
```

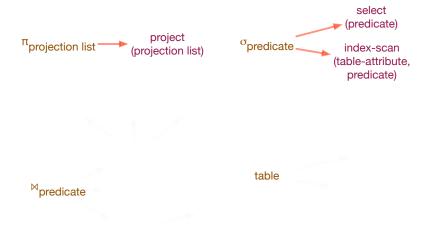
```
(student \bowtie_{student.cid=course.cid}
\sigma_{course.name='ADBS'} (course))
```

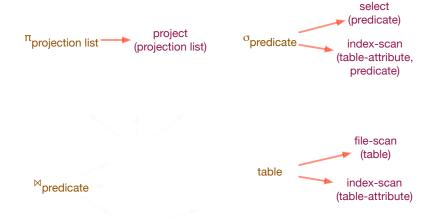
#### Algebraic operations

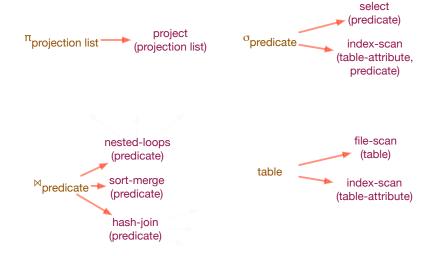
- $\pi_{student.id}$ , student.name
- $\bullet$   $\bowtie_{student.cid} = course.cid$
- $\sigma_{course.name} = `ADBS'$

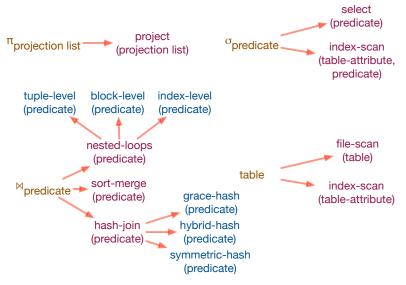






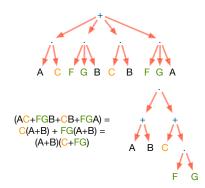






## Math analogy

- 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

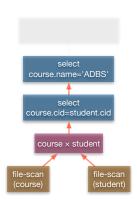


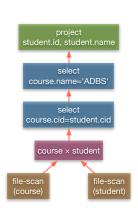


file-scan (course)

(student)



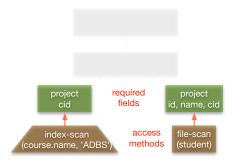




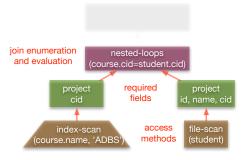
### SQL query



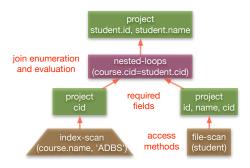
### SQL query



```
SQL query
```



```
SQL query
```



#### 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



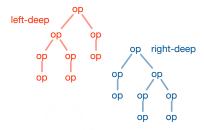
### A note on duplicates

- 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

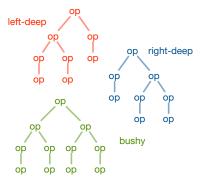
- There are two types of plan, according to their shape
  - ► Deep (left or right)
  - ► Bushy
- Different shapes for different objectives



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- There are two types of plan, according to their shape
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- 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



### Outline

#### 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

## 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

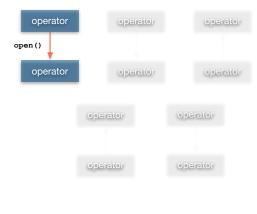
## What happens in practice

- 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*

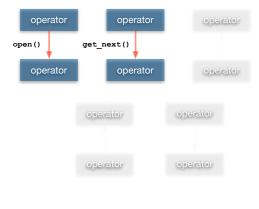
- Also known as a cursor
- Three basic calls
  - ▶ open()
  - ▶ get\_next()
  - ▶ close()
- Have you ever accessed a database through external code?
  - For example: exec sql declare cursor in embedded SQL, ResultSets in Java/JDBC, etc.



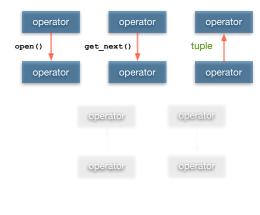
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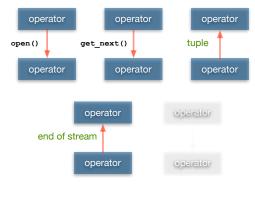
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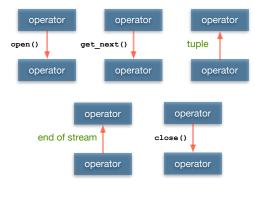
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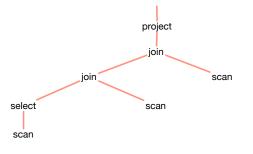


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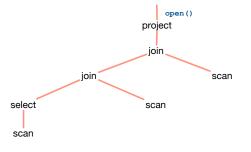


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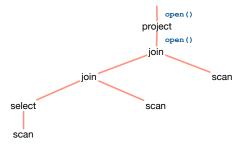




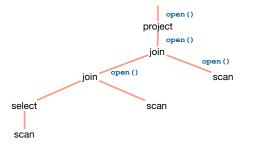
- All calls are propagated downstream
- The query engine makes the calls to the topmost operator only



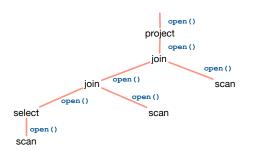
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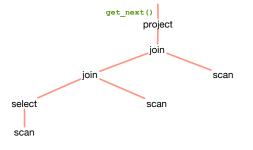
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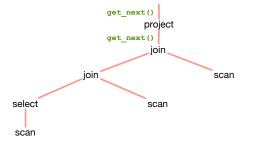
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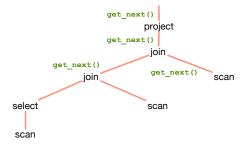


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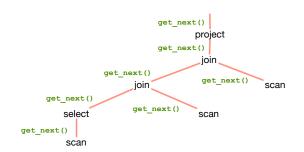
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# Call propagation



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- The query engine makes the calls to the topmost operator only

#### 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

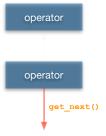
- The *iterator interface* is what *operators* use to *communicate*
- But how it is implemented, can be entirely different
  - ► The *reason* is that there might be *need for buffering*
  - ► Three possibilities
    - ★ Push model (buffering in the operator making the calls)
    - ★ Pull model (buffering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

- 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()

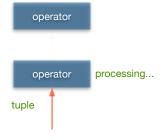
operator

operator

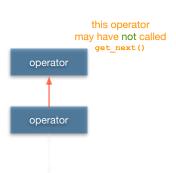
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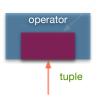
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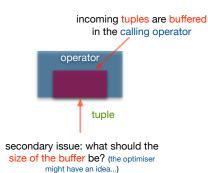
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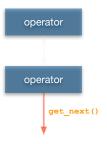




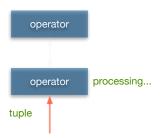




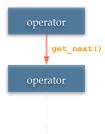
- The *inverse* of the *push model*
- If the lower operator is done processing a tuple it does not propagate it
  - ▶ It waits until the operator above it makes a get\_next() call



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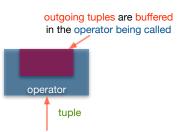
propagation only after a get next() call is made



- 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







outgoing tuples are buffered in the operator being called



same question: what should the size of the buffer be? (again, the optimiser might have an idea...)

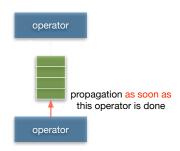
- The *connections* become *first-class citizens*
- Streams are queues of tuples connecting the operators
- Propagations and get\_next() calls are synchronised on each stream



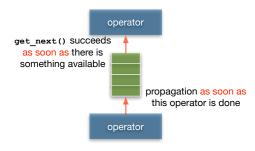
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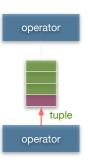
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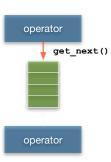
- This time, there is no question!
- When the lower operator is done, it propagates the tuple
- When the top operator is ready, it calls get\_next() on the incoming stream



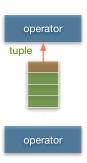
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## Why all this?

- 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

#### Outline

#### 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<sub>R</sub> and pages P<sub>R</sub>
  - 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

Tuple-level nested loops for each tuple  $r \in R$  do

### Tuple-level nested loops

for each tuple  $r \in R$  do for each tuple  $s \in S$  do

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• R is the outer relation

# Tuple-level nested loops

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for each tuple r \in R do
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```

- 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
  - - \* 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
  - $ightharpoonup cost(R \bowtie S) = P_R + P_R \cdot P_S$
  - ▶ 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

### 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

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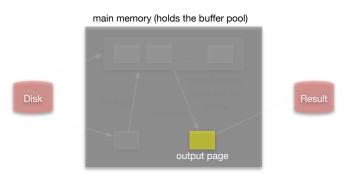
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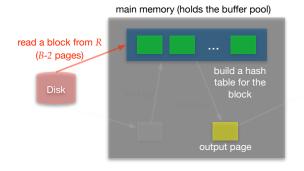
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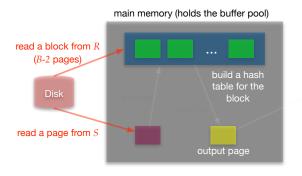
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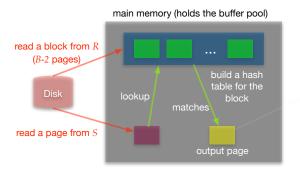




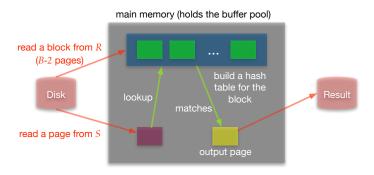








Result



#### How much does it cost?

- The *outer relation* is *still scanned once* (*P*<sub>R</sub> pages)
- The inner relation is scanned  $\lceil \frac{P_R}{B-2} \rceil$  times
  - ► Each scan costs P<sub>S</sub> 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

#### 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

### Index nested loops

Assumption: there is an index on S.b

for each tuple  $r \in R$  do

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#### Index nested loops

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```
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
  - $ightharpoonup cost(R \bowtie S) = T_R + T_R \cdot f \cdot T_S \cdot avg\_lookup$

### 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

- 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

	1*
Ī	2*
	2*

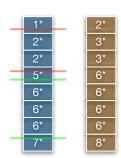




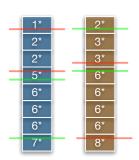




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while (more tuples in inputs) do {
while (r.a < gs.b) do advance r
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while (more tuples in inputs) do {

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while (r.a > gs.b) do advance gs // a group might begin here
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add \langle r, s \rangle to the result; advance s  // produce result
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```

#### What is the cost?

- We know the *cost* of *externally sorting* either *relation*:  $2 \cdot P_R \cdot \log P_R$ , or  $2 \cdot P_S \cdot \log P_S$
- The merge phase is essentially one scan of each sorted input: P<sub>R</sub> or P<sub>S</sub> (these scans are always clustered)
- $\bullet \ cost(R \bowtie S) = P_R \cdot (2 \cdot \log P_R + 1) + P_S \cdot (2 \cdot \log P_S + 1)$ 
  - ▶ 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 <u>large groups</u> in the <u>two relations</u>, then we <u>may</u> have to <u>do</u> 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<sub>i</sub> load it in memory, scan S<sub>i</sub> 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  $P_i$ ,  $i \in [1, m]$ 

## The simple algorithm

#### Simple hash join

```
Assumption: m partitions, each partition P_i fits in main memory
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```
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  $P_i$ 

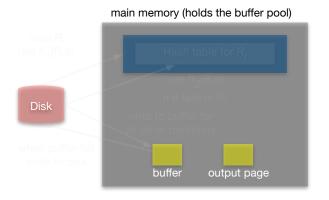
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## The simple algorithm

#### Simple hash join

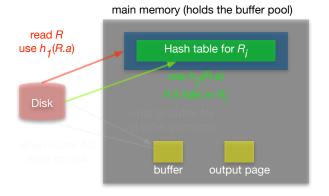
```
Assumption: m partitions, each partition P_i fits in main memory for all partitions P_i, i \in [1, m] for each r \in R read r and apply hash function h_1(r.a) if r falls into P_i apply hash function h_2(r.a) and put it in an in-memory hash table for P_i otherwise, write it back out to disk for each 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



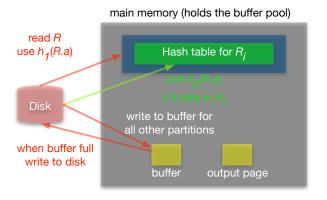
Result

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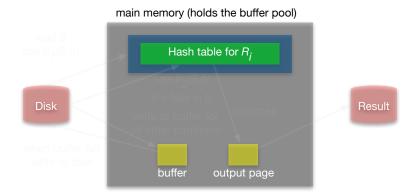


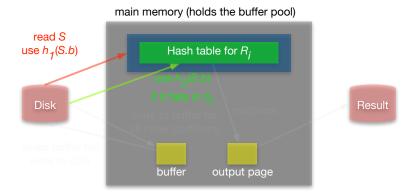


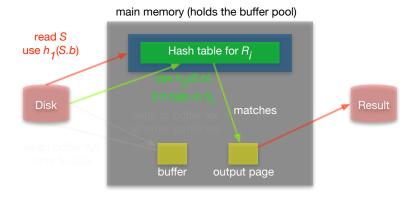
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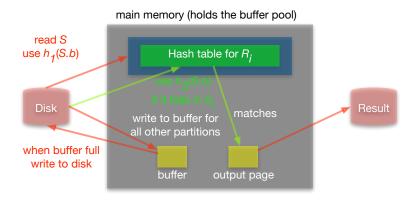












#### What is the cost?

- Assume equal partition sizes, input T, P<sub>T</sub> 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!

### Grace hash join

for each  $r \in R$  read r and add it to the buffer page for  $h_1(r.a)$ 

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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 {
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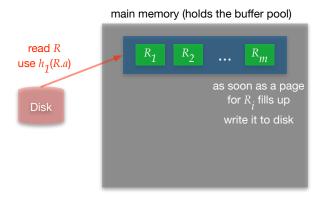
## How it works — partitioning R



#### main memory (holds the buffer pool)



## How it works — partitioning R



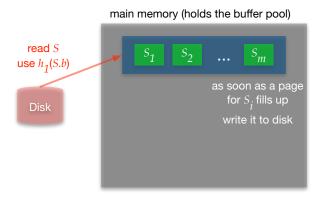
# How it works — partitioning *S*

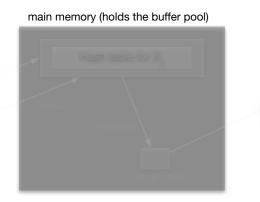


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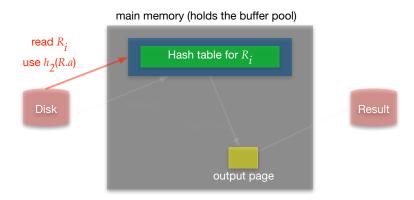


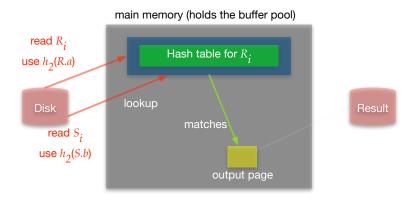
# How it works — partitioning *S*

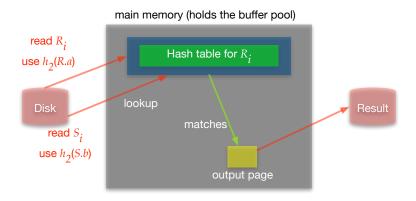




Result







#### 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<sub>S</sub>
- $cost(R \bowtie S) = 3 \cdot (P_R + P_S)$ 
  - ▶ 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

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- Size of each partition is  $\lceil \frac{P_R}{B-1} \rceil$
- Size of the hash table is  $\lceil \frac{f \cdot P_R}{B-1} \rceil$  (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

- 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  $\lceil \frac{P_R}{B-1} \rceil$
- Size of the hash table is  $\lceil \frac{f \cdot P_R}{B-1} \rceil$  ( $f = fudge \ factor$  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,  $B > \left\lceil \frac{f \cdot P_R}{R-1} \right\rceil + 2 \Rightarrow B > \sqrt{f \cdot P_R}$



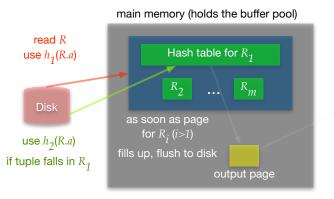
## Hybrid hash join

- 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

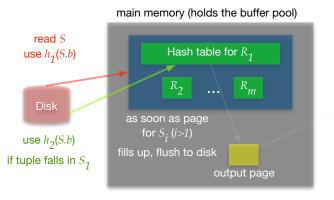
#### How it works

- 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 R<sub>1</sub> 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

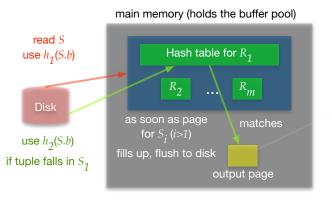




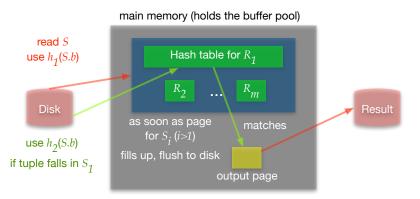












## Savings over grace hash join

- 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
  - ▶ 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.)

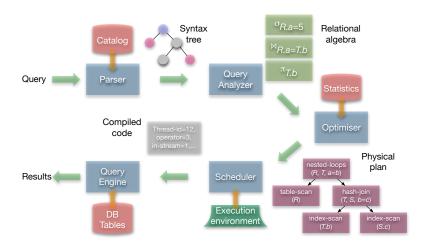
- Iteration-based methods
  - Essentially, nested loops
  - Very simple to implement, but if implemented poorly very inefficient
  - ▶ But also *very useful* because they *evaluate non-equi-join predicates*
- Order-based methods
  - Sort the inputs, merge them afterwards
  - ► Well-behaved cost 3-4 passes over the data will do the trick

# Summary (cont.)

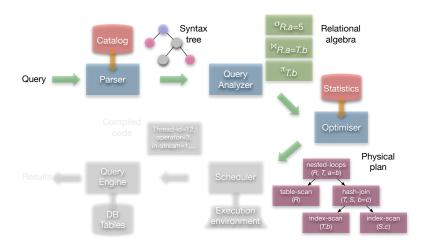
- Partition-based methods
  - Simple hash join, Grace hash join, and hybrid hash join
  - ▶ If there is extra memory, hybrid hash join's behaviour is excellent
- Figuring out the best join algorithm for a particular pair of inputs is the job of the query optimiser
- Which, along with good implementations,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

#### Outline

## Query cycle



# 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

#### 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 2<sup>5</sup> ⋅ 5! = 3840
  - ▶ If we add one extra join algorithm, the number of possible plans becomes  $2^4 \cdot 2^5 \cdot 5! = 61440$

## 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

# 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

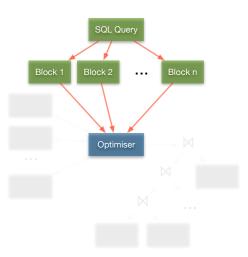
- Mapping SQL queries to relational algebra
  - Query blocks, uncorrelated vs. correlated queries
- Optimisation of a single query block
- Equivalence rules
- Statistics and cardinality estimation
- Search space exploration
  - Dynamic programming (System-R)

## Outline

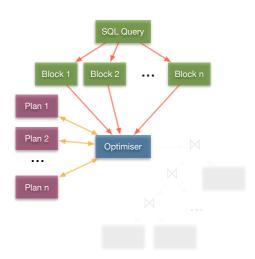
- SQL queries are optimised by decomposing them into a collection of query blocks
- A block is optimised in isolation, resulting in a plan for a block
- Plans for blocks are combined to form the complete plan for the query



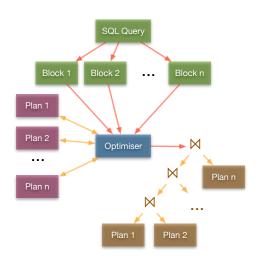
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### What is a block?

- 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)

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### 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
    s.rating = ( select max(s2.rating) from sailors s2)
group by    s.sid
having    count** > 1
```

```
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
                     s.rating = ( )
            group by s.sid
                     count(*) > 1
            having
         outer
        block
select
        s.sid, min(r.day)
from
        sailors s, reserves r, boats b
where
        s.sid = r.sid and r.bid = b.bid and
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                     count(*) > 1
         outer
                                            select max(s2.rating)
         block
                                            from
                                                   sailors s2
select
        s.sid, min(r.day)
from
         sailors s, reserves r, boats b
                                                   nested
where
         s.sid = r.sid and r.bid = b.bid and
                                                    block
         b.color = 'red' and
         s.rating = (select max(s2.rating)
                     from
                            sailors s2)
group by s.sid
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```

```
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            where
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            group by s.sid
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                                             reference
         outer
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                                             from
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group by s.sid
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```

#### 

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

## Relational algebra

```
\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))))
```

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- More specifically, extended relational algebra

## Relational algebra — before

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```

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

## Relational algebra — before

 $\pi_{s,sid,min(r,dav)}$ 

```
\begin{split} & \textit{having}_{\textit{count}(*)>2}(\\ & \textit{group by}_{\textit{s.sid}}(\\ & \sigma_{\textit{s.sid}-r.sid \land r.bid=b.bid \land b.color=red \land s.rating=nested-value}(\\ & \textit{sailors} \times \textit{reserves} \times \textit{boats})))) \end{split}
```

## Relational algebra — after

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```

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

- 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

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- Commutativity

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

# Cartesian products and joins

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  - $ightharpoonup R \times S \equiv S \times R$
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- Their combination

- Selection-projection commutativity
  - $\bullet$   $\pi_a$  (  $\sigma_c$  (R))  $\equiv \sigma_c$  (  $\pi_a$  (R))
  - ▶ iff every attribute in c is included in the set of attributes a

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  - $ightharpoonup \sigma_c (R \times S) \equiv R \bowtie_c S$
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  - ▶ iff the attributes in c appear only in R and not in S
- Selection distribution/replacement

  - iff  $c_1$  is relevant only to R and  $c_2$  is relevant only to S

# Among operations (cont.)

- Projection-Cartesian product commutativity
  - $\quad \bullet \quad \pi_a \ (R \times S) \equiv \pi_{a_1}(R) \times \pi_{a_2}(R)$
  - ▶ iff a₁ is the subset of attributes in a appearing in R and a₂ is the subset of attributes in a appearing in S

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  - ▶ iff same as before and every attribute in c appears in a
- Attribute elimination

  - iff  $a_1$  subset of attributes in R appearing in either a or c and  $a_2$  is the subset of attributes in S appearing in either a or c

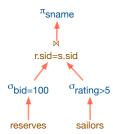
#### What do we have and what do we need?

- 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*

### Outline

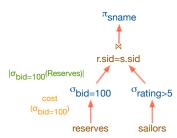
#### Cost estimation

- 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



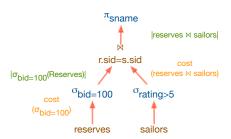
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### Cost and cardinality

- 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

```
SQL query  \begin{array}{ll} \text{select} & \textbf{a}_1, \textbf{a}_2, \dots \textbf{a}_k \\ \text{from} & R_1, R_2, \dots R_n \\ \text{where} & P_1 \text{ and } P_2 \text{ and } \dots \text{ and } P_m \end{array}
```

#### SQL query

 $\begin{array}{ll} \text{select} & \textbf{\textit{a}}_1, \textbf{\textit{a}}_2, \dots \textbf{\textit{a}}_k \\ \text{from} & R_1, R_2, \dots R_n \end{array}$ 

where  $P_1$  and  $P_2$  and ... and  $P_m$ 

# Maximum output cardinality

 $|R_1|\cdot |R_2|\cdot\ldots\cdot |R_n|$ 

#### SQL query

select  $a_1, a_2, \dots a_k$ 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

$$f_{P_1} \cdot f_{P_2} \cdot \ldots \cdot f_{P_m}$$

#### SQL query

select  $a_1, a_2, \dots a_k$ from  $R_1, R_2, \dots R_n$ 

where  $P_1$  and  $P_2$  and ... 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

# 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 column<sub>1</sub> has a matching value in column<sub>2</sub>; given a value in column<sub>1</sub>, the predicate is just a selection
  - ► Again, an approximation

•  $column > value \rightarrow \frac{(high(column) - value)}{(high(column) - low(column))}$ 

```
• column > value \rightarrow \frac{(high(column) - value)}{(high(column) - low(column))}
```

```
 \bullet \ \ \textit{value}_1 < \textit{column} < \textit{value}_2 \rightarrow \frac{(\textit{value}2 - \textit{value}1)}{(\textit{high}(\textit{column}) - \textit{low}(\textit{column}))}
```

- ullet column > value  $\to \frac{(\mathit{high}(\mathit{column}) \mathit{value})}{(\mathit{high}(\mathit{column}) \mathit{low}(\mathit{column}))}$
- $value_1 < column < value_2 \rightarrow \frac{(value_2 value_1)}{(high(column) low(column))}$
- column in list  $\rightarrow$  number of items in list times s.f. of column = value

- $column > value \rightarrow \frac{(high(column) value)}{(high(column) low(column))}$
- $value_1 < column < value_2 \rightarrow \frac{(value_2 value_1)}{(high(column) low(column))}$
- 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

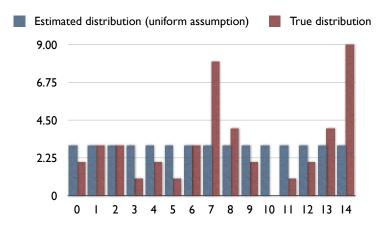
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- $\bullet \ \ P_1 \lor P_2 \to f_{P_1} + f_{P_2} f_{P_1} \cdot f_{P_2}$

## 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



## 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

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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

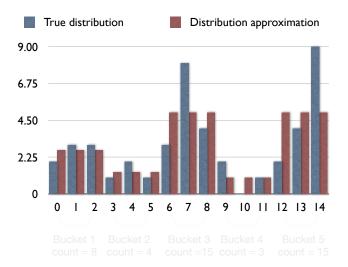
- Elegant data structures to capture value distributions
  - Not affected by the uniform distribution assumption (though this is not entirely true)
- They offer *trade-offs* between *size* and *accuracy* 
  - ► The more memory that is dedicated to a histogram, the more accurate it is
  - ▶ But also,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

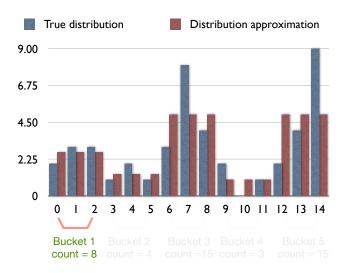
## Desirable histogram properties

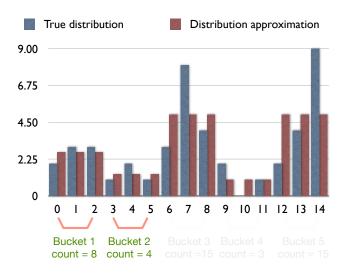
- Small
  - ► Typically, a DBMS will allocate a *single page* for a histogram!
- Accurate
  - ► Typically, less than *5% error*
- Fast access
  - ► Single lookup access and simple algorithms

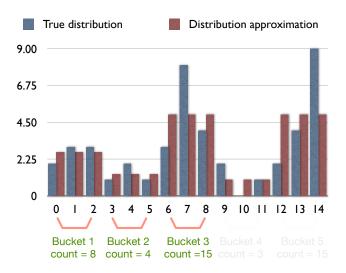
## Mathematical properties

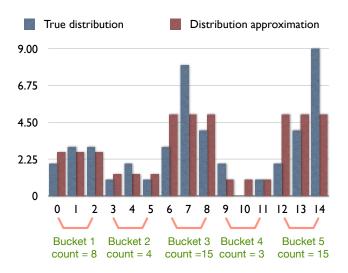
- A histogram approximates the value distribution for attribute X of table T
- The value distribution is partitioned into a number of b subsets, called buckets
- There is a partitioning constraint that identifies how the partitioning takes place
  - Different constraints, lead to different classes of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



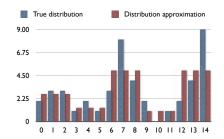






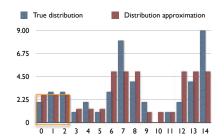


- The *total range* is *divided* into *sub-ranges* of *equal width*
- Each *sub-range* becomes a *bucket*
- The total number of tuples in each bucket is stored



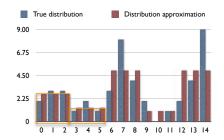


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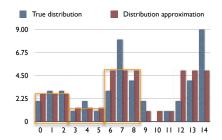


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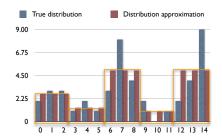
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## Equi-width histogram construction

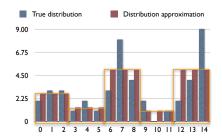
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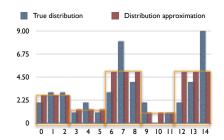
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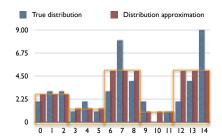
min	max	count
0	2	8
3	5	4
6	8	15
9	11	3
12	14	15

- To estimate the output cardinality of a range query
  - ► The *starting bucket* is *identified*



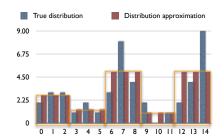
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- To estimate the output cardinality of a range query
  - ► The *starting bucket* is *identified*
  - ► The histogram is then scanned forward until the ending bucket is identified



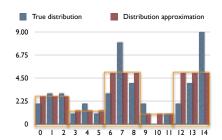
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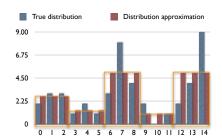
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  - ► Within each bucket the uniform distribution assumption is made

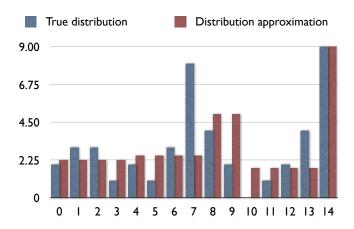


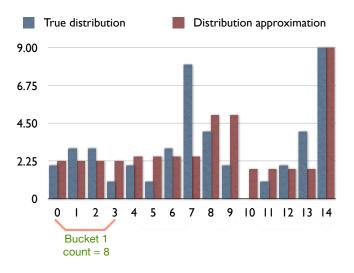
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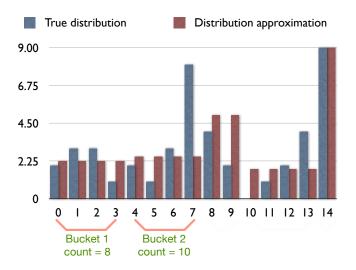
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  - Within each bucket the uniform distribution assumption is made
- $6 \le v \le 10$ :  $\frac{3}{3} \cdot 15 + \frac{2}{3} \cdot 3 = 17$

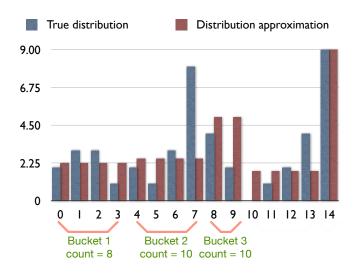


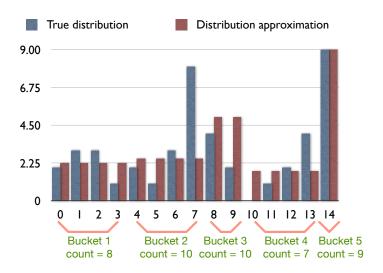
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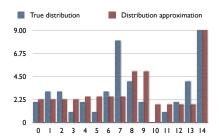






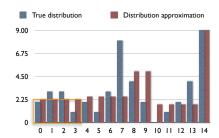


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- The same schema as in equi-width histograms is used
- In fact, the *same algorithm* is used for *estimation* (!)
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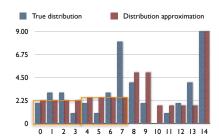
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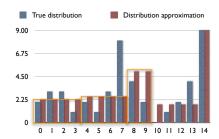
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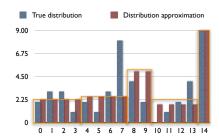
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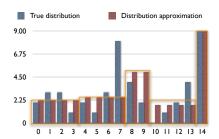
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#### Comparison

- 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

## Outline



# 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

- 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

• *Identify* the *cheapest* way to *access every* single *relation* in the query, *applying local predicates* 

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- 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







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

local predicates

```
select name, title, salary, dname
from emp, dept, job
where job.title='Clerk' and
    dept.location = 'Edinburgh' and
    emp.dno = dept.dno and
    emp.job = job.job
```

in predicates interesting orders

# emp name dno job salary Smith 50 12 8500 Jones 50 5 15000 Doe 51 5 9500

#### dept

dno	dname	location
50	MFG	Edinburgh
51	Billing	London
52	Shipping	Glasgow

job

JOD	
job	title
5	clerk
6	typist
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```

join predicates

#### emp

name	dno	job	salary
Smith	50	12	8500
Jones	50	5	15000
Doe	51	5	9500

#### dept

dno	dname	location
50	MFG	Edinburgh
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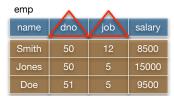
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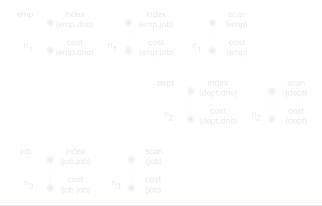
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#### local predicates

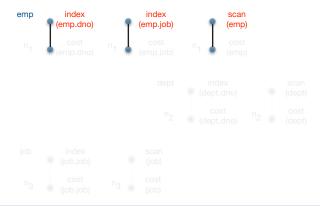
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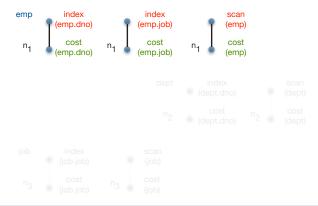
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# Access methods and local predicates



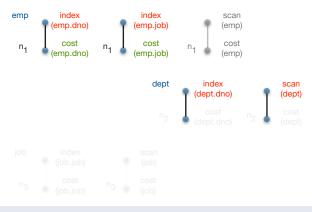
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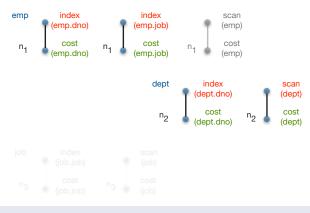




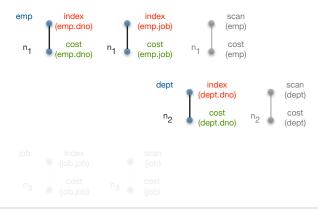
• Scanning emp is the most expensive method for emp; emp.dno and emp.job are interesting orders



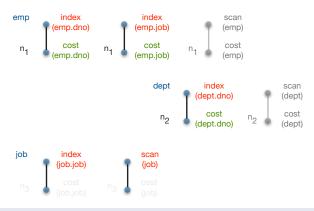
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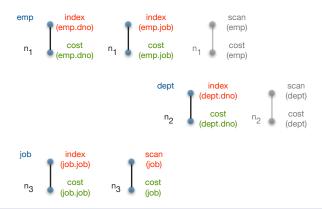
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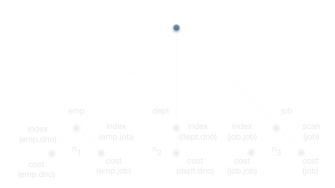
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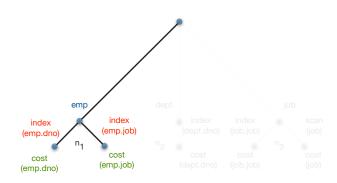


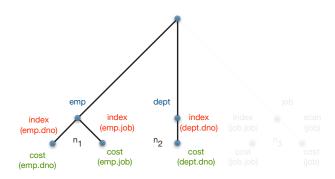
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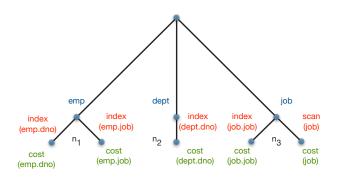


- Scanning emp is the most expensive method for emp; emp.dno and emp.job are interesting orders
- Scanning dept is the most expensive method for dept; dept.dno is an interesting order
- Scanning job is the cheapest method for job; but, job.job is an interesting order





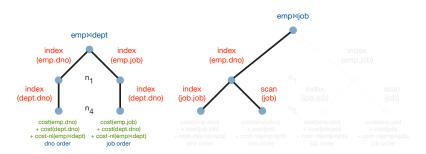












■ Both emp 

dept results are in different interesting orders so they are propagated.

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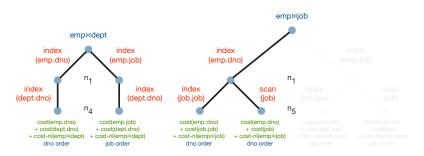
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dept 

dept

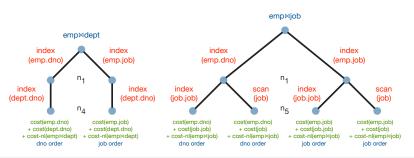


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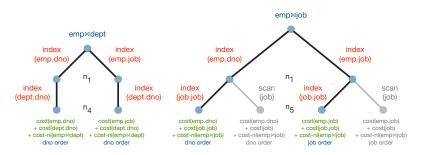
dept results are in different interesting orders so they are propagated.

Output

Description:



● Both emp ⋈ dept results are in different interesting orders so they are propagated



- Both emp ⋈ 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



•  $cost(emp \bowtie dept) \neq cost(dept \bowtie emp)$  so we will enumerate dept's joins even though we have an alternative for generating the same result (same for  $job \bowtie emp)$ 



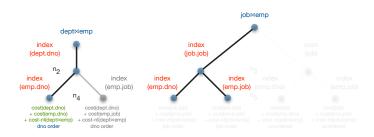
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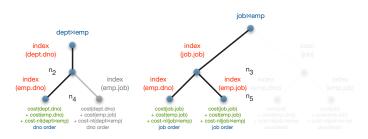
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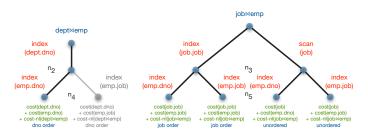
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- Since there is no dept ⋈ job predicate in the query, that join is not enumerated (same for job ⋈ dept)



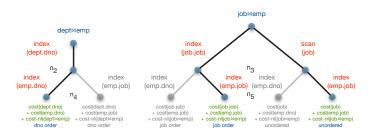
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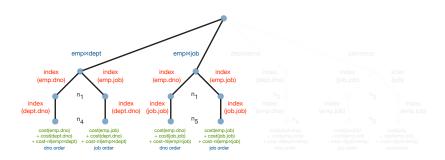
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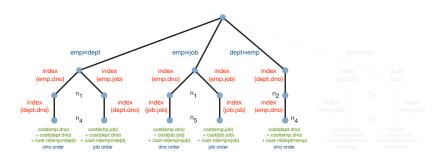


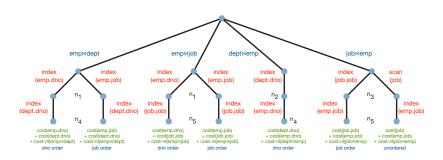
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- Since there is no dept ⋈ job predicate in the query, that join is not enumerated (same for job ⋈ dept)
- The unordered result for job ⋈ emp is propagated because it is the cheapest overall



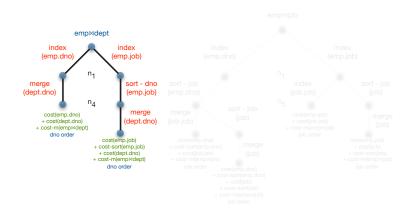




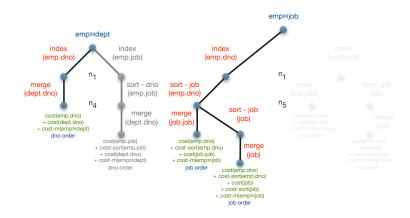


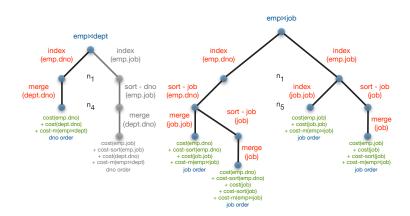


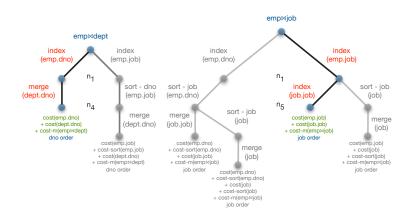






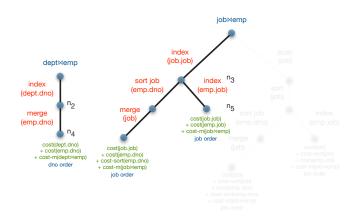


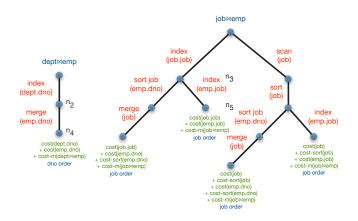


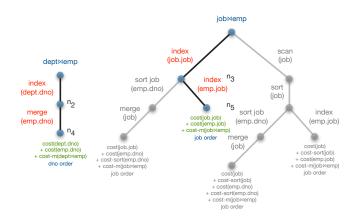




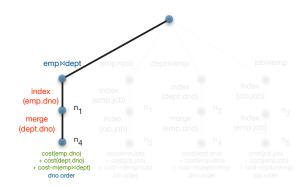


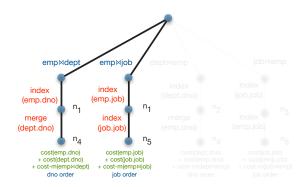


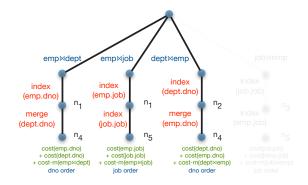


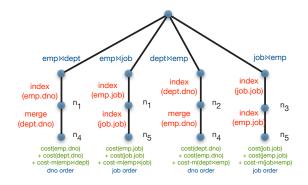








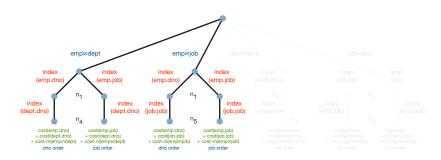




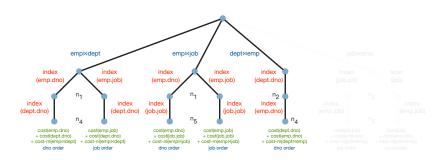




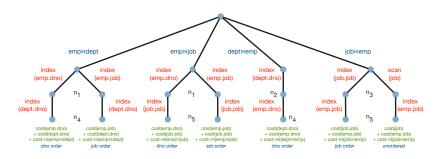
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- For each pair or relations, for each different join order and for each interesting order for that pair one plan is propagated
- An unordered result is only propagated if it is the cheapest overall for a pair in a given join order

#### Three relations

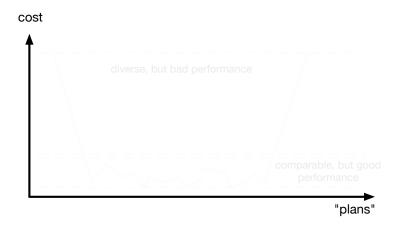
- Repeat the process
  - ► For *every pair* of two relations
  - ► For *every join* method
  - ► For *every access method* of the *remaining relation*
  - ► Find the cheapest way to join the third relation with the pair
    - **★** Estimate cardinalities
    - ★ Estimate the cost of computing the join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

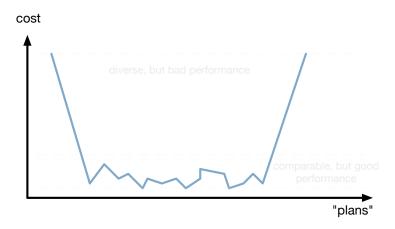
### Rule-based optimisation

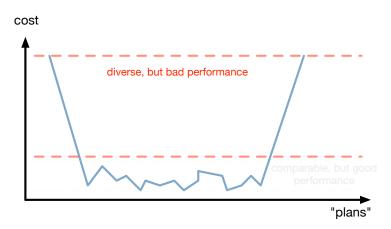
- Basically an issue of if-then rules
  - If (condition list) then apply some transformation to the plan constructed so far
    - \* Estimate the cost of the new plan, keep it only if it is cheaper than the original
  - ► The *order* in which the *rules are applied* is *significant*
  - ► As a *consequence*, 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

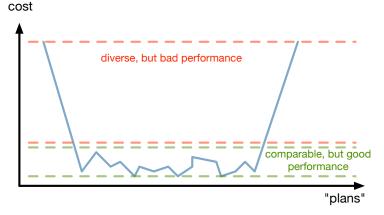
- Mostly useful in big queries (more than 15 joins or so)
- The *problem* is one of *exploring a bigger portion* of the search space
  - ► So, every once in a while the optimiser "jumps" to some other part of the search space with some probability
- As a consequence, it gets to explore parts of the search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



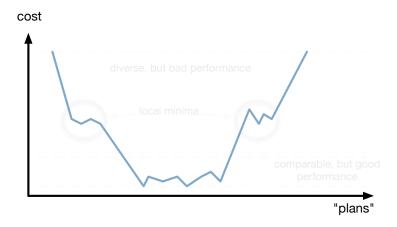


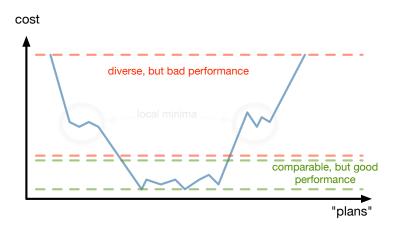


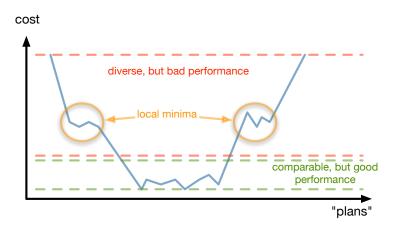












### Final step — the entire plan

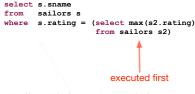
- The *optimiser* has produced *plans* for *each query block*
- The *question* is now one of *combining* the *sub-plans* to *formulate* the *entire query plan*
- The strategy used depends on whether the outer and nested queries are correlated or not
  - ► If they are, then in all probability the two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

### Uncorrelated queries

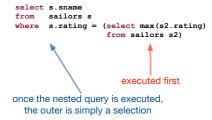
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once the nested query is executed, the outer is simply a selection

### Uncorrelated queries

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- The nested query feeds the outer query with results



- 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

```
select s.sname
       sailors s
       exists (select *
               where r.bid = 103 and
                      s.sid = r.sid)
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                        σr.bid=103
                         reserves
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       sailors
                         5r.bid=103
                          reserves
                       this could be
                       an entire plan
```

### 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!

# Outline

# 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*

# Outline

## 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 (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

# Outline

### **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!

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- *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

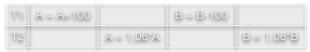
## T1

Begin A = A+100 B = B-100 End

T2

Begin A = 1.06\*A B = 1.06\*B End

### Acceptable schedule



### Problematic schedule



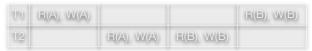


### Acceptable schedule

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

### Problematic schedule



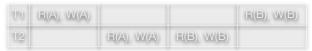


### Acceptable schedule

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	



### Acceptable schedule

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

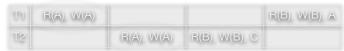
### Problematic schedule

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

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

# Scheduling

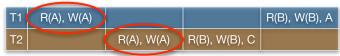
- A schedule is a sequence of reads and writes for some transaction workload incorporating all actions of the workload's transactions
  - ► Serial schedule: the actions of different transactions are not interleaved
  - ► Equivalent schedules: for any database state, the effect of executing the first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







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



Unrepeatable reads (RW conflicts)





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

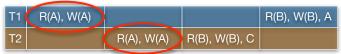


Unrepeatable reads (RW conflicts)

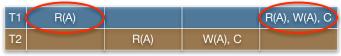




### Reading uncommitted data (WR conflicts, or "dirty reads")

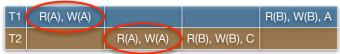


Unrepeatable reads (RW conflicts)

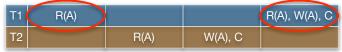


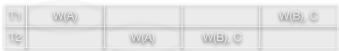


### Reading uncommitted data (WR conflicts, or "dirty reads")

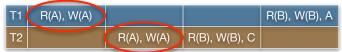


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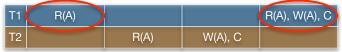


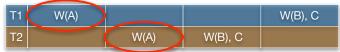


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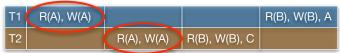


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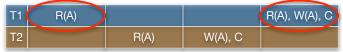


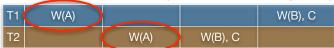


### Reading uncommitted data (WR conflicts, or "dirty reads")



### Unrepeatable reads (RW conflicts)





 Before a transaction "touches" a DB object it has to obtain a lock for it

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  - ► S (Shared) lock for reading

- 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

- Before a transaction "touches" a DB object it has to obtain a lock for it
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- Strict two-phase locking (Strict 2PL)

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  - ► 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

#### 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

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- Before a transaction "touches" a DB object it has to obtain a lock for it
  - S (Shared) lock for reading
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  - ► 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*
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  - ► 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

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  - ▶ 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<sub>i</sub> is aborted, then all its actions have to be undone; not only that, but if T<sub>j</sub> reads an object written by T<sub>i</sub>, T<sub>j</sub> needs to be aborted as well (cascading aborts)
- Most systems avoid cascading aborts with the following rule:
  - ▶ If  $T_i$  writes an object  $T_i$  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

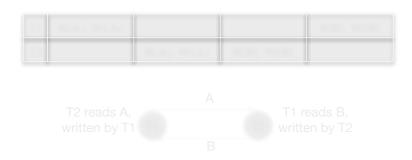
### Outline

# 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



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

T2 reads A, T1 reads B, written by T1 Written by T2

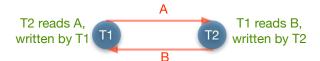
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T1	R(A), W(A)			R(B), W(B)
T2		R(A), W(A)	R(B), W(B)	



#### 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)
  - reads before it starts reading it and an X (eXclusive) lock for everything it writes before it starts writing

► Each transaction must obtain an S (Shared) lock for everything it

- ► 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  $T_j$  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

- Create a waits-for graph
  - ► *Nodes* are *transactions*
  - ► There is an edge from T<sub>i</sub> to T<sub>j</sub> if T<sub>i</sub> is waiting for T<sub>i</sub> 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)		
ТЗ						S(C)	R(C)			X(A)
T4									X(B)	

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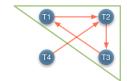
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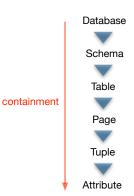
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# Multiple granularity locks

- 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

#### held lock

wanted lock

				014 100	••		
		NL	IS	IX	SIX	S	Х
	NL	Υ	Υ	Υ	Υ	Υ	Y
5	IS	Υ	Υ	Υ	Υ	Υ	N
אמוונסט	IX	Υ	Υ	Υ	N	N	N
<b>8</b>	SIX	Υ	Υ	N	N	N	N
	S	Υ	Υ	N	N	Υ	N
	Х	Υ	N	N	N	N	N

#### 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

• T1 scans R, and updates a few tuples

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- T3 reads all of R
  - ► T3 gets an S lock on the entire relation

# 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)

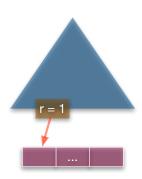
- If we *relax* the *assumption* that the *DB* is a *fixed collection* of objects, *even Strict 2PL* will *not assure serialisability*!
  - ► T1 locks all pages containing sailor records with rating = 1, and finds oldest sailor (say, age = 71)
  - ► Next, *T2 inserts* a *new sailor*: rating = 1, age = 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

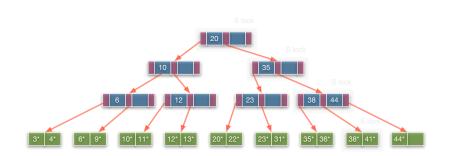
- How can we efficiently lock a particular node?
  - ► This is *entirely different* than *multiple granularity locking* (why?)
- One solution: ignore the tree structure, just lock pages while traversing the tree, following 2PL
  - ► Terrible performance
  - ► Root node (and many higher level nodes) become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

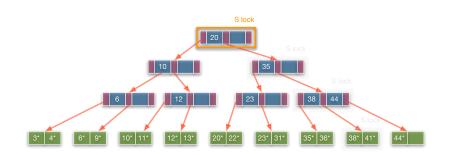
# Key observations

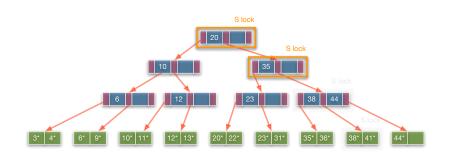
- Higher levels of the tree only direct searches to leaf pages
- For insertions, a node on a path from the root to a modified leaf must be locked (in X mode, of course), only if a split can propagate up to it from the modified leaf (similar point holds for deletions)
- We can exploit these observations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

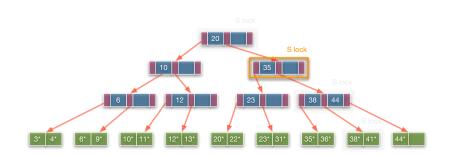
# The basic algorithm

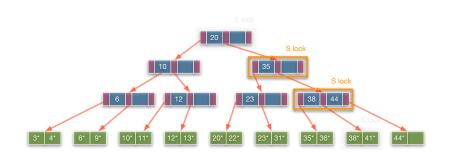
- Search: start at root and descend; repeatedly, S lock child then unlock parent
- Insert/Delete: start at root and descend, obtaining X locks as needed; once child is locked, check if it is safe:
  - Safe node: a node such that changes will not propagate up beyond this node
    - \* Insertion: node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

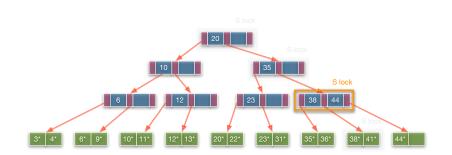


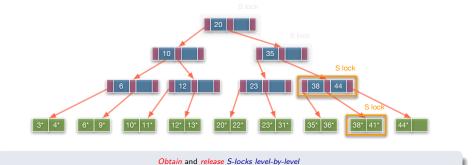


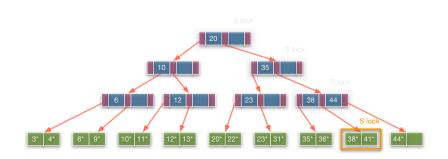


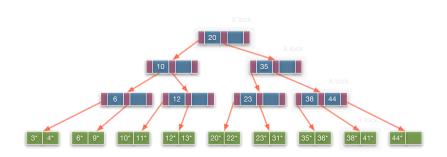


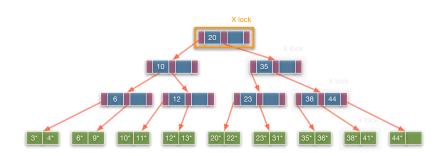


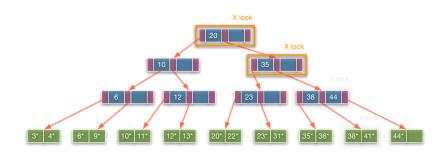


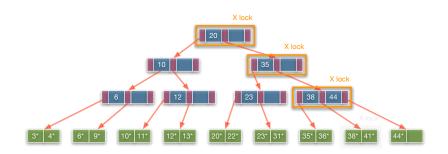


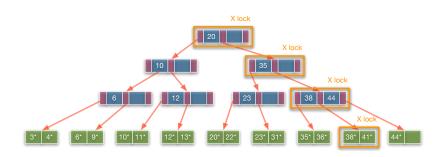


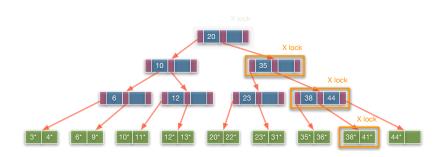


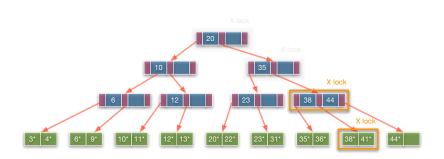


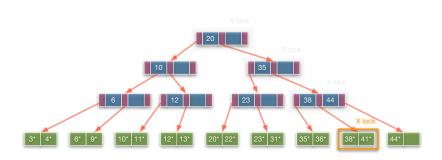


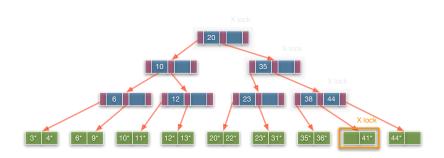


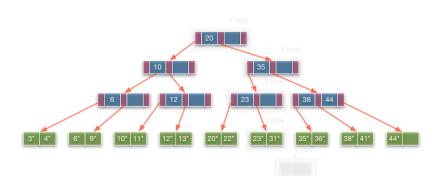


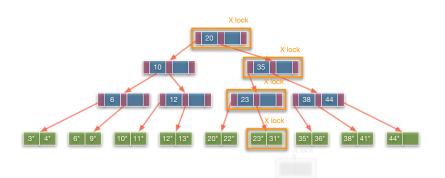


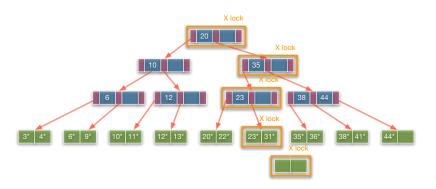


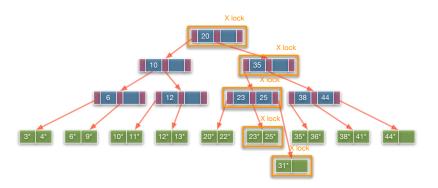


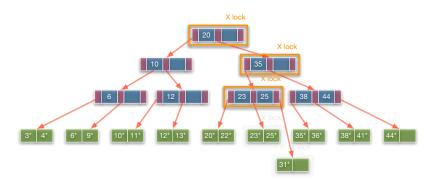


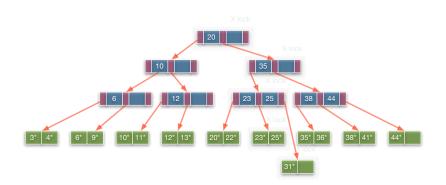






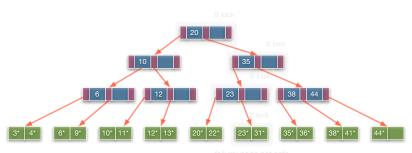




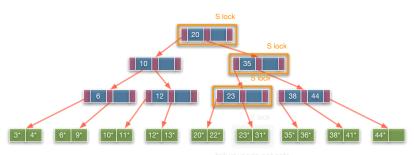


# Optimistic B+tree locking

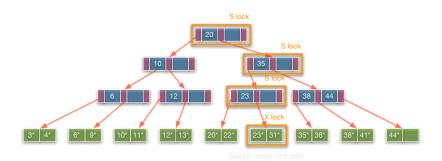
- Search: as before
- Insert/delete: set locks as if for search, get to the leaf, and set X lock on the leaf
  - If the leaf is not safe, release all locks, and restart transaction, using previous insert/delete protocol
- "Gambles" that only leaf node will be modified; if not, S locks set on the first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

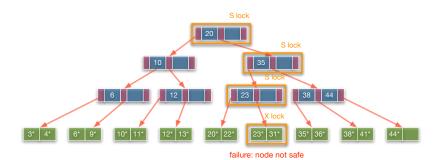


tallure: node not sate



tallure: node not sate





### 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



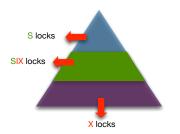
- The likelihood that we will need an X lock decreases as we move up the tree
- Set S locks at high levels, SIX locks at middle levels, X locks at low levels

# Hybrid approach

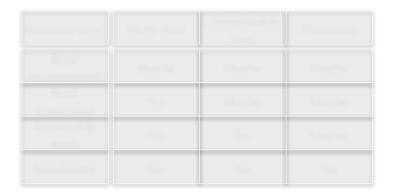


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Isolation level	Dirty read	Unrepeatable read	Phantoms
Read uncommitted		Selection (	Selection (
Read committed	-	Salary San	Marylan
Repeatable reads	-	-	bilanyilay
Serialisable			

Isolation level	Dirty read	Unrepeatable read	Phantoms
Read uncommitted	Maybe	Maybe	Maybe
Read committed	-	Salary Sa	bilayday
Repeatable reads	-	-	bilaying
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Isolation level	Dirty read	Unrepeatable read	Phantoms
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Serialisable	No	No	No

#### Outline

#### 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

#### Review: ACID properties

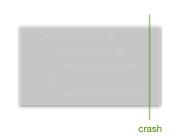
• Atomicity: all the actions in a transaction are executed as a single atomic operation; either they are all carried out or none are

• *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?

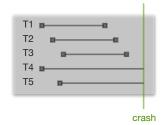
- Atomicity
  - Transactions may abort; their effects need to be undone
- Durability
  - What if the system stops running?



Transactional semantics

# What can go wrong?

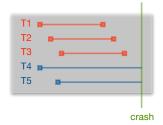
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Transactional semantics

# What can go wrong?

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



#### Transactional semantics

- T1, T2, T3 should be durable
- T4, T5 should be aborted

#### 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

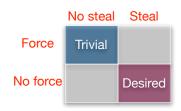
# More on the buffer pool

- 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



# More on the buffer pool

- Two issues: force and steal
- Force: when a data page is modified it is written straight to disk
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  - ► But durable
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  - ► 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)

### 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

# Write-ahead logging

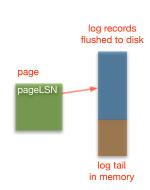
- The log adheres to the write-ahead protocol (WAL)
  - Must force the log record for an update before the corresponding data page gets to disk
  - Must force all log records for a transaction before it commits
- #1 guarantees atomicity
- #2 guarantees durability

#### 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 <u>log record</u> has a unique <u>log</u> 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 flushed! SN
  - ► The max LSN flushed so far
- WAL: before a page is written, pageLSN < flushedLSN</li>



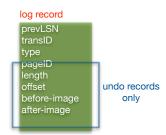
# Log records

- 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

# log record prevLSN transID type pageID length offset before-image after-image

## Log records

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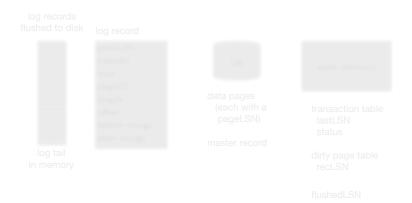


### Other log-related state

- Transaction table: one entry per active transaction
  - Contains transaction id, status (running/committed/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

## 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)



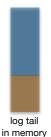
### log records flushed to disk log record prevLSN length offset before-image log tail





in memory

log records flushed to disk



log record

prevLSN transID type pageID length offset before-image after-image DB

data pages (each with a pageLSN)

master record

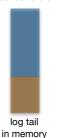
-----

ransaction table lastLSN status

dirty page table recLSN

flushedLSN

log records flushed to disk



log record

prevLSN transID type pageID length offset before-image after-image DB

data pages (each with a pageLSN)

master record

main memory

transaction table lastLSN status

dirty page table recLSN

flushedLSN

#### 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 *lastLSN* 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

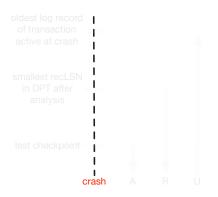
#### Transaction commit

- 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

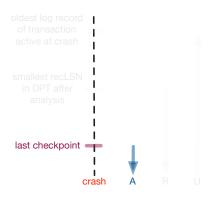
# Recovery: big picture



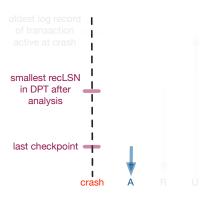
- Start from a checkpoint (found via master record)
- Three phases
  - Analysis: figure out which transactions committed since the checkpoint, and which failed
  - ► REDO all actions
    - ★ Repeat history
  - UNDO effects of failed transactions



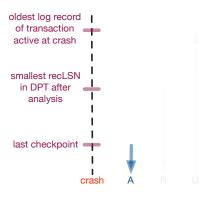
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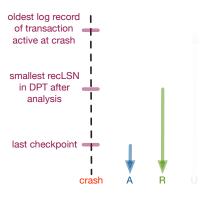
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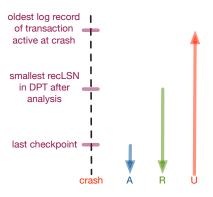
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### Additional issues

- 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

### Outline

## 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

- 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

- 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

- 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

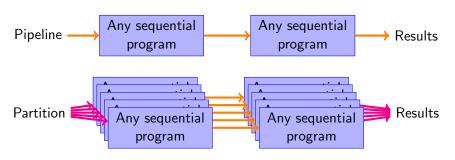
- Use *checkpoints* to *periodically record consistent states* and *limit* the amount of the *log* that needs to be *scanned during recovery*
- Recovery in three phases
  - ► Analysis: from checkpoint, figure out REDO and UNDO extents
  - ► REDO: repeat entire history
  - ► UNDO: delete effects of failed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



Partitioning: *split* inputs, *merge* outputs

## 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*

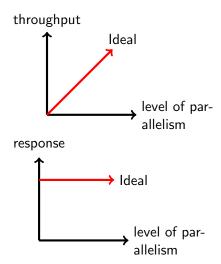
# Terminology

### Speed-up

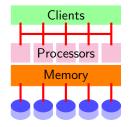
More resources means proportionally less time for given amount of data (throughput)

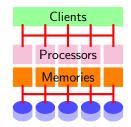
### Scale-up

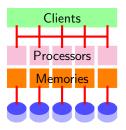
If resources increased in proportion to increase in data size, time is constant



### 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

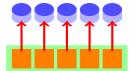
### Shared nothing

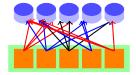
- Hard to program
- Cheap to build
- Easy and ideal to speed/scale up

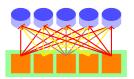
# Different types of parallelism

- 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

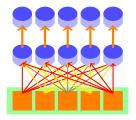
- Indifferent for equi-joins
- Range-queries complicated
- Load-balanced

### 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()

  - 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

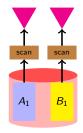
- During the *first phase*, *partitions* are *distributed* to *different sites* 
  - A good hash function automatically distributes work evenly
- Second phase is local at each site
  - ► Almost *always* the *winner* for *equi-join*
- Good use of split/merge makes it easier to build parallel versions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

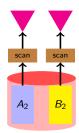
## Dataflow network for parallel join



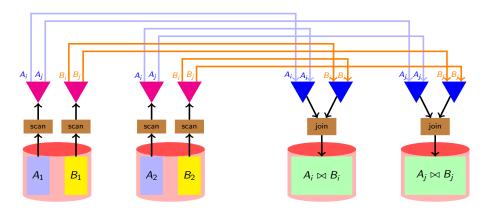


## Dataflow network for parallel join



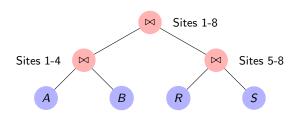


## Dataflow network for parallel join



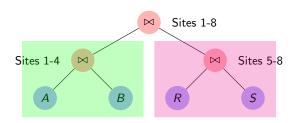
## Complex parallel query plans

- 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



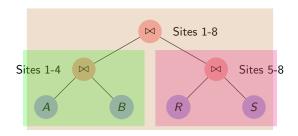
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### Observations

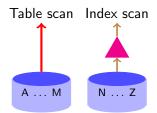
- It is *relatively easy* to build a *fast parallel query executor*
- It is hard to write a robust and world-class parallel query optimizer
  - ► There are many *tricks*
  - ► One quickly hits the *complexity barrier*
  - ► Still open research

## 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 ≠ 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"</p>
  - ▶ *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*