L21 Query Execution & Optimization Continued

SELECT a FROM R

$$\pi_a(R)$$

 π_a I

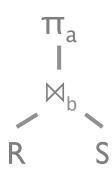
SELECT a FROM R WHERE a > 10

$$\pi_a(\sigma_{a>10}(R))$$

$$\begin{array}{c} \pi_a \\ I \\ \sigma_{a>10} \\ I \\ R \end{array}$$

SELECT a
FROM R JOIN S
ON R.b = S.b

$$\pi_a(\bowtie_b(R,S))$$



Push vs Pull?

Push

Operators are input-driven

As operator (say reading input table) gets data, push it to parent operator.

Pull

Operators are demand-driven

If parent says "give me next result", then do the work

Are cursors push or pull?

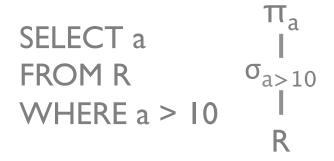
Naïve execution (operator at a time)

read R

filter a>10 and write out

read and project a

Cost: B + M + M

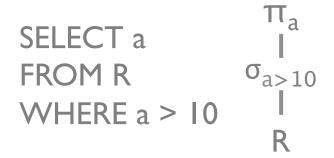


- B # data pages
- M # pages matched inWHERE clause

Could we do better?

```
Pipelined exec (tuple/page at a time) read first page of R, pass to \sigma filter a > 10 and pass to \pi project a (all operators run concurrently) Cost: B
```

Note: can't pipeline some operators! e.g., sort, some joins, aggregates why?



B # data pagesM # pages matched inWHERE clause

What if R is indexed?

Hash index

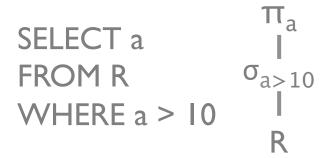
Not appropriate

B+Tree index

use a>10 to find initial data page

scan leaf data pages

Cost: log_FB + M



- B # data pages
- M # pages matched inWHERE clause

Access Paths

Choice of how to access input data is called the Access Path

file scan or

index + matching condition (e.g., a > 10)

Access Paths

Sequential Scan doesn't accept any matching conditions

Hash index search key < a,b,c> accepts conjunction of equality conditions on *all* search keys e.g., a=1 and b=5 and c=5 will (a=1) and b=5) work? why?

Tree index search key <a,b,c>
accepts conjunction of terms of *prefix* of search keys
typically best with equality on all but last column

```
e.g., a = 1 and b = 5 and c < 5 will (a = 1 \text{ and } b > 5) work? will (a > 1 \text{ and } c > 9) work?
```

How to pick Access Paths?

Selectivity

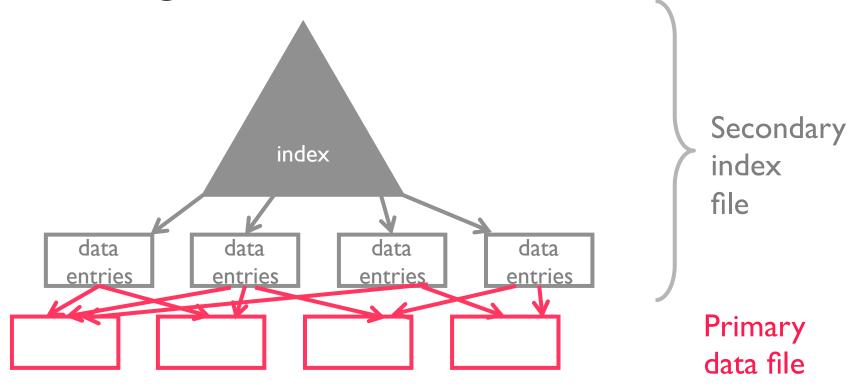
ratio of # outputs satisfying predicates vs # inputs
0.01 means I output tuple for every 100 input tuples

Assume

attribute selectivity is independent

```
if selectivity(a=1) = 0.1, selectivity(b>3) = 0.6
selectivity(a=1 and b>3) = 0.1*0.6 = 0.06
```

High level index structure



What is a data entry?

actual data record

<search key value, rid>

<search key value, rid_list>

How to pick Access Paths?

Hash index on <a, b, c>a = |, b = |, c = | how to estimate selectivity?

- pre-compute attribute statistics by scanning data
 e.g., a has 100 values, b has 200 values, c has 1 value
 selectivity = 1 / (100 * 200 * 1)
- 2. How many distinct values does hash index have? e.g., 1000 distinct values in hash index
- make a number up "default estimate" is the fancy term

System Catalog Keeps Statistics

```
System R
```

```
NCARD "relation cardinality" # tuples in relation
```

```
TCARD # pages relation occupies
```

```
ICARD # keys (distinct values) in index
```

```
NINDX pages occupied by index
```

min and max keys in indexes

Statistics were expensive in 1979!

Catalog stored in relations too!

What Optimization Options Do We Have?

Access Path Predicate push-down
Join implementation
Join ordering

In general, depends on operator implementations. So let's take a look

Predicate Push Down

SELECT a
$$\Pi_a$$
 $\sigma_{a>10}$ Π_a $\sigma_{a>10}$ Π_a $\sigma_{a>10}$ Π_a Π_a

Which is faster if B+ Tree index: (a) or (b)?

- (a) $log_F(B) + M pages$
- (b) B pages

It's a Good Idea, especially when we look at Joins

Projection with DISTINCT clause

need to deduplicate e.g., π_{rating} Sailors

Two basic approaches

Sort: fundamental database operation sort on rating, remove dups on scan of sorted data Hash:

partition into N buckets remove duplicates on insert

Index on projected fields scan the index pages, avoid reading data

The Join

Core database operation join of 10s of tables common in enterprise apps

Join algorithms is a large area of research

e.g., distributed, temporal, geographic, multi-dim, range, sensors, graphs, etc

Discuss three basic joins nested loops, indexed nested loops, hash join

Best join implementation depends on the query, the data, the indices, hardware, etc

Nested Loops Join:

```
# outer ⋈₁ inner
# outer JOIN inner ON outer.1 = inner.1
for row in outer:
    for irow in inner:
        if row[0] == irow[0]: # could be any check
            yield (row, irow)
```

Very flexible

Equality check can be replaced with any condition Incremental algorithm

Cost: M + MN

Is this the same as a cross product?

Nested Loops Join

What this means in terms of disk IO

tableA join tableB; tableA is "outer"; tableB is "inner" M pages in tableA, N pages in tableB, T tuples per page

 $M + T \times M \times N$

for each tuple t in tableA, (M pages,TM tuples) scan through each page pi in the inner (N pages) compare all the tuples in pi with t

Nested Loops Join: Order?

Does order matter?

$$M + T \times M \times N$$

$$N + T \times N \times M$$

Scan "outer" once; Scan "inner" multiple times: If inner is small IO cost is M + N!

Indexed Nested Loops Join

```
for row in outer:
   for irow in index.get(row[0], []):
     yield (row, irow)
```

Slightly less flexible

Only supports conditions that the index supports

Indexed Nested Loops Join

What this means in terms of disk IO

outer join inner on sid

M pages in outer, N pages in inner, T tuples/page inner has primary key index on sid

Cost of looking up in index is C₁

predicate on outer has 5% selectivity

```
M + T \times M \times 0.05 \times C_1
```

for each tuple t in the outer: (M pages, TM tuples) if predicate(t): (5% of tuples satisfy pred) lookup_in_index(t.sid) (C_I disk IO)