# Lecture 10 10/6/1014

**Query Optimization** 

PS 2 due Wednesday.

#### Selinger

Famous paper. Pat Selinger was one of the early System R researchers; still active today.

Lays the foundation for modern query optimization. Some things are weak but have since been improved upon.

Idea behind query optimization:

```
(Find query plan of minimum cost )
```

How to do this?

(Need a way to measure cost of a plan (a cost model) )

#### single table operations

```
how do i compute the cost of a particular predicate?
compute it's "selectivity" - fraction F of tuples it passes
```

how does Selinger define these? -- based on type of predicate and available statistics

what statistics does system R keep?

- NCARD "relation cardinality" -- number of tuples in relation
- TCARD # pages relation occupies
- ICARD keys (distinct values) in index
- NINDX pages occupied by index
- min and max keys in indexes

(have to realize that the complexity of statistics you could keep in 1978 was pretty simple!)

# Estimating selectivity F:

```
col = val
    F = 1/ICARD() (if index available)
    F = 1/10 (where does this come from?)

col > val
    high key - value / high key - low key (if index available)
    1/3 o.w.

col1 = col2
    1/MAX(ICARD(col1, col2))
    1/10 o.w.

ex: suppose emp has 1000 records, dept has 10 records
total records is 1000 * 10, selectivity is 1/1000, so 10 tuples expected to pass join
(note that this is wrong if doing key/fk join on emp.did = dept.did, which will produce 1000 results!)
```

Note that selectivity is defined relative to size of cross product for joins!

p1 and p2

```
p1 or p2
```

```
1 - (1-F1) * (1-F2)
```

then, compute access cost for scanning the relation. how is this defined?

(in terms of number of pages read)

equal predicate with unique index: 1 [btree lookup] + 1 [heapfile lookup] + W

(W is CPU cost per predicate eval in terms of fraction of a time to read a page)

range scan:

```
clustered index, boolean factors: F(preds) * (NINDX + TCARD) + W*(tuples read)
```

unclustered index, boolean factors: F(preds) \* (NINDX + NCARD) + W\*(tuples read) unless all pages fit in buffer -- why?

seq (segment) scan: TCARD + W\*(NCARD)

Is an index always better than a segment scan? (no)

#### multi-table operations

how do i compute the cost of a particular join?

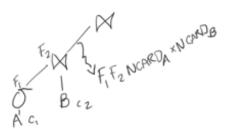
algorithms:

NL(A,B,pred)Cost(A) + NCARD(A) \* Cost(B)

Note that inner is always a relation; cost to access depends on access methods for B; e.g., w/ index -- 1 + 1 + W w/out index -- TCARD(B) + W\*NCARD(B)

Cost(A) is cost of subtree under outer

How to estimate # NCARD(outer)? product of F factors of children, cardinalities of children example:



Merge\_Join\_x(P,A,B), equality pred

Cost(A) + Cost(B) + sort cost

(Saw cost models for these last time)

At time of paper, didn't believe hashing was a good idea

Overall plan cost is just sum of costs of all access methods and join operators Then, need a way to enumerate plans Iterate over plans, pick one of minimum cost

### Problem:

```
Huge number of plans. Example:
suppose I am joining three relations, A, B, C
Can order them as:
(AB)C
A(BC)
(AC)B
A(CB)
(BA)C
B(AC)
(BC)A
B(AC)
(CA)B
C(AB)
(CB)A
C(BA)
Is C(AB) different from (CA)B?
Is (AB)C different from C(AB)?
     yes, inner vs. outer
n! strings * # of parenthetizations
How many parenthetizations are there?
Consider N=1,2,3,4:
  A: (A)
  AB: ((A)(B))
  ABC: ((AB)C), (A(BC))
  ABCD:(((AB)C)D), ((A(BC))D), ((AB)(CD)), (A((BC)D)), (A(B(CD)))
The numbers of plans for N=1,2,3,4 are:
  plans(1) = 1
  plans(2) = 1
  plans(3) = 2
  plans(4) = 5
Generally, plans(N) = choose(2(N-1),(N-1))/(N) *
* The Art of Computer Programming, Volume 4A, page 440-450
==> n! * choose(2(N-1),(N-1))/(N)!
6 * 2 == 12 for 3 relations
(study break -- postgres)
Ok, so what does Selinger do?
```

Push down selections and projections to leaves

Now left with a bunch of joins to order.

Selinger simplifies using 2 heuristics? What are they?

```
- only left deep; e.g., ABCD \Rightarrow (((AB)C)D) show
```

- ignore cross products

e.g., if A and B don't have a join predicate, doing consider joining them

still n! orderings. can we just enumerate all of them?

```
10! -- 3million
20! -- 2.4 * 10 ^ 18
```

so how do we get around this?

Estimate cost by dynamic programming:

idea: if I compute join (ABC)DE -- I can find the best way to combine ABC and then consider all the ways to combine that with DE.

I can remember the best way to compute (ABC), and then I don't have to re-evaluate it. Best way to do ABC may be ACB, BCA, etc -- doesn't matter for purposes of this decision.

algorithm: compute optimal way to generate every sub-join of size 1, size 2, ... n (in that order).

```
R <--- set of relations to join
for ∂ in {1...IRI}:
     for S in {all length ∂ subsets of R}:
           optioin(S) = a join (S-a), where a is the single relation that minimizes:
                 cost(optioin(S-a)) +
                 min cost to join (S-a) to a +
                 min. access cost for a
example: ABCD
only look at NL join for this example
A = best way to access A (e.g., sequential scan, or predicate pushdown into index...)
                          " B
B = " "
C = "
                          " C
D = "
                          " D
\{A,B\} = AB \text{ or } BA
\{A,C\} = AC \text{ or } CA
\{B,C\} = BC \text{ or } CB
\{A,D\}
\{B,D\}
{C,D}
\{A,B,C\} = remove A - compare A(\{B,C\}) to (\{B,C\})A
           remove B - compare ({A,C})B to B({A,C})
```

remove C - compare C({A,B}) to ({A,B})C

{A,C,D} {A,B,D} {B,C,D}

```
remove A - compare A({B,C,D}) to ({B,C,D})A
{A,B,C,D} =
               remove B
               remove C
                remove D
Complexity:
number of subsets of size 1 * work per subset = W+
number of subsets of size 2 * W +
number of subsets of size n * W+
n + n + n ... n
1 2 3 n
number of subsets of set of size n = power set of n = 2^n
(string of length n, 0 if element is in, 1 if it is out; clearly, 2<sup>n</sup> such strings)
(reduced an n! problem to a 2^n problem)
what's W? (at most n)
so actual cost is: 2<sup>n</sup> * n
```

# So what's the deal with sort orders? Why do we keep interesting sort orders?

Selinger says: although there may be a 'best' way to compute ABC, there may also be ways that produce interesting orderings -- e.g., that make later joins cheaper or that avoid final sorts.

So we need to keep best way to compute ABC for different possible sort orders.

so we multiply by "k" -- the number of interesting orders

#### how are things different in the real world?

- real optimizers consider bushy plans (why?)

n=12 --> 49K vs 479M

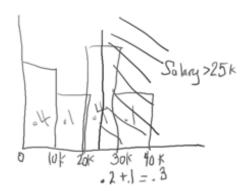
- selectivity estimation is much more complicated than selinger says and is very important.

# how does selinger estimate the size of a join?

- selinger just uses rough heuristics for equality and range predicates.
- what can go wrong?
   consider ABCD
   suppose sel (A join B) = .1
   everything else is .01
   If I don't leave A join B until last, I'm off by a factor of 10

- how can we do a better job? (multi-d) histograms, sampling, etc.

example: 1d hist



example: 2d hist

