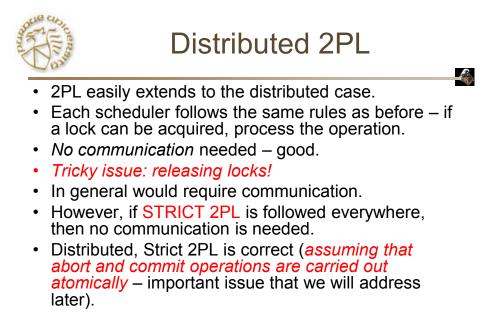




2 Phase Locking

- 1. To grant a lock, the scheduler *checks if a conflicting lock* has already been assigned, if so, *delay*, otherwise *set lock and grant* it.
- A lock cannot be released at least until the DM acknowledges that the operation has been performed.
- 3. Once the scheduler releases a lock for a txn, it may not subsequently acquire any more locks (on any item) for that txn.



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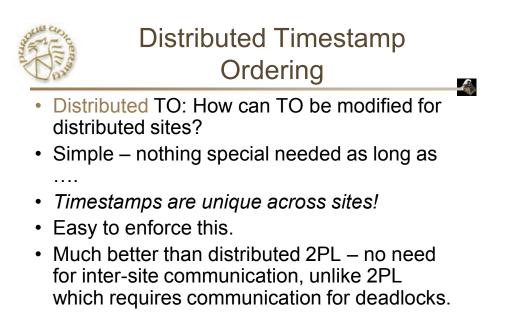


Distributed Deadlocks

- As with centralized 2PL, distributed 2PL suffers from deadlocks. Moreover, these can be distributed deadlocks! E.g. if x and y are at different sites.
- Solutions:
 - Timeouts
 - Deadlock Detection
 - Deadlock Prevention
- Timeouts are easy local decision, but may be overreacting.



- The TM assigns each txn, T_i, a unique timestamp, ts(Ti).
- No two txns share a timestamp.
- A TO scheduler enforces:
- TO Rule: if p_i[x] and q_j[x] are conflicting operations, then the DM processes p_i[x] before q_j[x] iff ts(T_i) < ts(T_j).

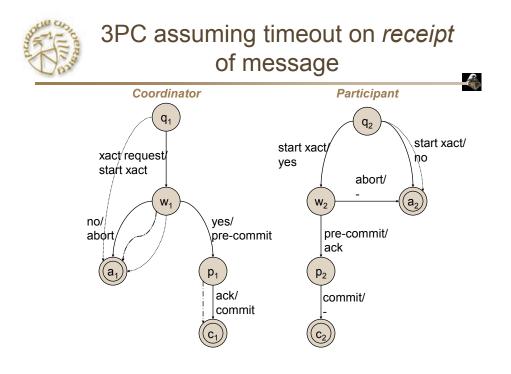


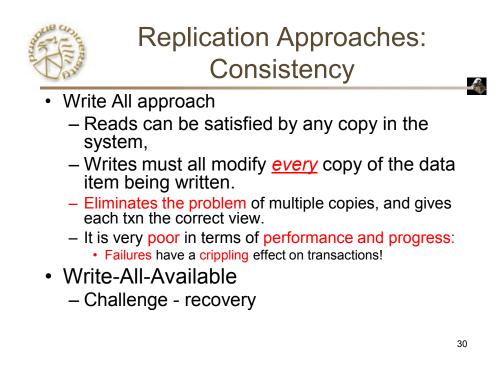


Recovery

- We will focus on system failures.
- Following the failure, the DBMS is restarted.
- At the start of recovery, the contents of *volatile storage are discarded.*
- The stable storage is potentially inconsistent
- A CONSISTENT database state corresponding to exactly the set of txns that had committed (as far as the DM is concerned) must be reconstructed, i.e. C(H).
- This reconstruction uses only data in stable storage – Stable DB and the LOG.

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1 Copy Serializability

- The correctness definition for replicated databases is therefore that it should behave as though all transactions are executed in a <u>serial</u> <u>manner on a single copy database</u>.
- This is the notion of one copy serializability, I.e. 1SR.
- The user must be given a one copy view of the database.
- · How is this achieved?
- Read-only is easy. For writes we must manage carefully!

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Distributed Design Issues

- Why fragment?
- How to fragment?
- How much to fragment?
- How to test correctness?
- How to allocate?
- Information requirements?

Correctness of fragmentation

- Completeness
 - Decomposition of Relation *R* into *R*₁, *R*₂, ...*R*_n is complete if and only if each data item in R can also be found in some *R*_i
- Reconstruction
 - If Relation R is decomposed into $R_1, R_2, ..., R_n$, then there should exist some operator, that *R* can be reconstructed from $R_1, ..., R_n$.
- Disjointness
 - If Relation *R* is decomposed into *R*₁, *R*₂, ...*R*_n, and data item *d* is in *R*_j, then *d* should not be in any other fragment *R*_k, *k* <>j.

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PHF-Information Requirements

- Application Information
 - <u>Simple predicates</u>: Given *R*[*A*₁, *A*₂, ..., *A*_n], a simple predicate *p_i* is:
 - P_i : $A_i \theta$ Value
 - where θ is a comparison operator, Value is from the domain of attribute A_i
 - <u>Minterm predicates</u>: Given *R* and $P_r = \{p_1, p_2, \dots, p_m\}$, define $M = \{m_1, m_2, \dots, m_n\}$ as

where
$$p_i^* = p_i$$
 or $NOT(p_i)$.
 $M = \{m_i \mid m_i = \bigwedge_{pj \in \Pr} p_j^*\}, 1 \le i \le z$

Primary Horizontal Frag.

Definition:

- Where $F_i = \sigma_{F_i}(R), 1 \le j \le w$ Where F_i is a selection formula, which is (preferably) a minterm predicate.
- Therefore,
 - A horizontal fragment, R_i of relation R consists of all the tuples of R which satisfy a minterm predicate $m_i \rightarrow$
 - Given a minterm of predicates *M*, there are as many horizontal fragments of relation R as there are minterm predicates
 - Set of horizontal fragments also referred to as minterm fragments.

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PHF - Algorithm

- GIVEN: A relation R, the set of simple predicates P_r
- OUTPUT: The set of fragments of $R = \{R_1, R_2\}$ \dots, R_{W} which obey the fragmentation rules.
- Preliminaries:
 - $-P_r$ should be complete
 - $-P_r$ should be minimal

PHF - Example

- Two candidate relations: PAY and PROJ.
- Fragmentation of relation PAY
 - Application: check the salary info and determine raise.
 - Employee records kept at two sites
 application run at two sites
 - Simple predicates
 - p₁ : SAL <= 30000
 - p₂ : SAL > 30000
 - $P_r = \{p_1, p_2\}$ which is complete and minimal $P_r = P_r$
 - Minterm predicates
 - *m*₁ : (SAL <= 30000)
 - m_2 : NOT(SAL <= 30000) = (SAL>30000)

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Fragmentation of **PROJ**

· Applications:

- Find the name and budget of projects given their loc.— issued at three sites
- Access project information according to budget
 - One site accesses <=200000 another accesses > 200000

Simple Predicates

- For application 1:
 - p₁: LOC = "Montreal"
 - p₂: LOC = "New York"
 - *p*₃ : LOC = "Paris"
- For application 2:
 - P₄: BUDGET <= 200000
 - P₅: BUDGET > 200000
- $-P_r = P_r' = \{p_1, p_2, p_3, p_4, p_5\}$

PHF Example

• Fragmentation of PROJ contd:

- Minterm fragments left after elimination
- *m*₁: (LOC = "Montreal") AND (BUDGET <=200000)
- *m*₂: (LOC = "Montreal") AND (BUDGET>200000)
- m_3 : (LOC = "New York") AND (BUDGET <= 200000)
- $-m_4$: (LOC = "New York") AND (BUDGET > 200000)
- *m*₅: (LOC = "Paris") AND (BUDGET <=200000)
- *m*₆: (LOC = "Paris") AND (BUDGET >200000)

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PHF -- Example

PROJ₁

PNO	PNAME	BUDGET	LOC
P1	Instr.	150000	Montreal

 $PROJ_4$

PNO	PNAME	BUDGET	LOC
Р3	CAD/CA M	250000	New York

 $PROJ_2$

PNO	PNAME	BUDGET	LOC
P2	Database Develop.	135000	New York

PROJ₆

PNO	PNAME	BUDGET	LOC
P4	Maint.	310000	Paris

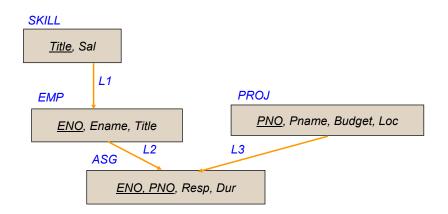
Derived Horizontal Fragmentation

- Defined on a member relation of a link according to a selection operation specified on its owner.
 - Each link is an equijoin
 - Equijoin can be implemented by means of semijoins.

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Derived Horizontal Fragmentation



VF – Information Requirements

Application Information

- Attribute affinities
 - A measure that indicates how closely related the attributes are
 - · This is obtained from more primitive usage data
- Attribute usage values
 - Given a set of queries $Q = \{q_1, q_2, ..., q_k\}$ that will run on the relation $R[A_1, A_2, ..., A_n]$,
 - Use(q_i, A_i) = 1 if A_i is referenced by qi, 0 otherwise
 - Use(qi,.) can be defined accordingly

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VF – Affinity Measure aff(Ai,Aj)

 The attribute affinity measure between two attributes Ai and Aj of a relation R with respect to the set of applications Q={q1, q2, ..., qk} is defined as follows:

$$aff(A_i, A_j) = \sum_{\text{all queries that accessAi and Aj}} (\text{query access})$$

query access=
$$\sum_{allsites} accessfreq of a query* \frac{access}{execution}$$

Bond Energy Algorithm

- Input: the AA matrix
- Output: the clustered affinity matrix CA (a perturbation of AA)
- 1. Initialization: Place and fix one of the columns of AA in CA
- 2. Iteration: Place the remaining n-I columns in the remaining I+1 positions in the CA matrix. For each column, chose the placement that makes the most contribution to the global affinity measure.
- 3. Row Order: Order the rows according to the columns.

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Selecting Alternatives

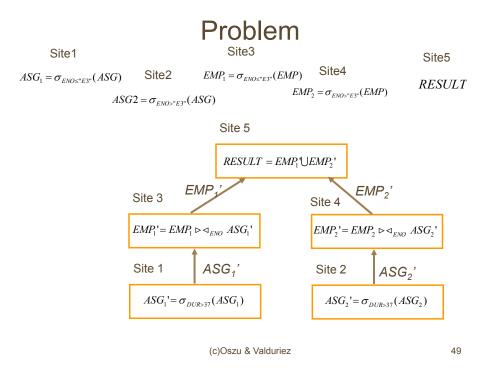
SELECT ENAME FROM EMP, ASG WHERE EMP.ENO = ASG.ENO AND DUR > 37.

Strategy 1:

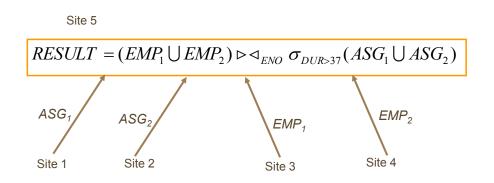
 $\prod_{ENAME} (\sigma_{EMP.ENO=ASG.ENO \land DUR>37}(EMP \times ASG))$

Strategy 2: $\prod_{ENAME}(EMP \triangleright \triangleleft_{ENO} (\sigma_{DUR>37}(ASG)))$

Strategy 2, avoids cartesian product.



Alternative 2



Cost of Alternatives

- Assume
 - Size(EMP) = 400; size(ASG)=1000
 - Tuple access cost (TAC) = 1unit; tuple xfer cost (TXC) =10units
- Strategy 1
 - Produce ASG': (10+10)*TAC = 20
 - Transfer ASG': (10+10)*TXC = 200
 - Produce EMP': (10+10)*TAC*2 = 40
 - Transfer EMP' to result site: (10+10)*TXC = 200
 - Total COST = 460.

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Cost of alternatives (cont)

- Strategy 2
 - Transfer EMP to site 5: 400*TXC = 4000
 - Transfer ASG to site 5: 1000*TXC = 10,000
 - Produce ASG': 1000*TAC = 1,000
 - Join EMP and ASG': 400*20*TAC = 8,000

- TOTAL COST = 23,000!!

Query Optimization Objectives

- Minimize a cost function

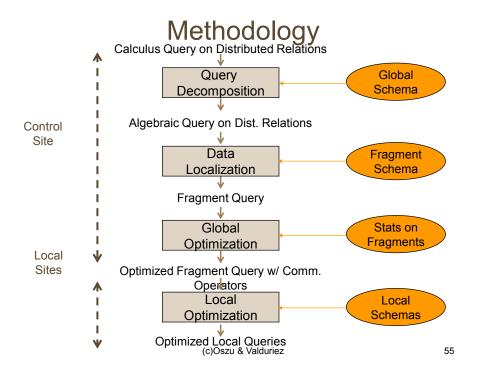
 I/O cost + CPU cost + communication cost
- These may have different weights in different distributed environments
- · Wide area networks
 - Communication cost will dominate
 - Low bandwidth
 - Low speed
 - High protocol overhead
 - Most algorithms ignore all other cost components

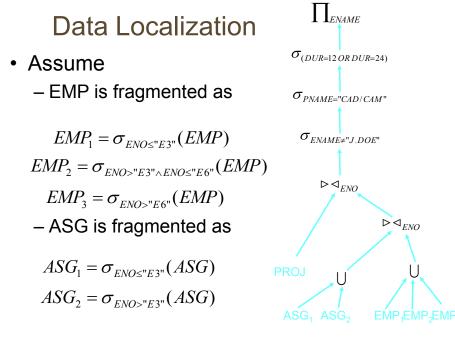
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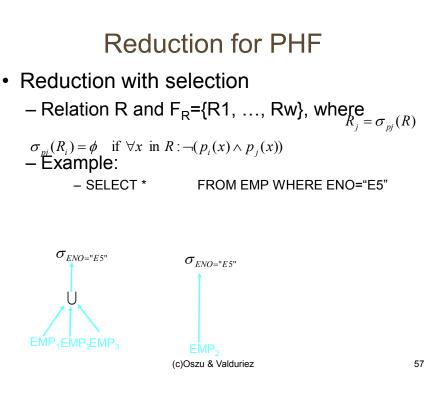
Complexity of Relational Operators

	Operation	Complexity
Assume Relations of cardinality <i>n</i> Sequential scan	Select, Project (without duplicate elimination)	O(n)
	Project (w/ duplicate elimination) Group	$O(n \log n)$
	Join Semijoin Division Set Operators	<i>O(n</i> log <i>n)</i>
	Cartesian Product (c)Oszu & Valduriez	O(n²)





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Reduction for PHF

- · Reduction with join
 - Possible if fragmentation is done on join attribute
 - Distribute join over unions $(R_1 \cup R_2) \triangleright \triangleleft S \Leftrightarrow (R_1 \triangleright \triangleleft S) \cup (R_2 \triangleright \triangleleft S)$
 - Given $R_i = \sigma_{pi}(R)$ and $R_j = \sigma_{pj}(R)$
 - $R_i \triangleright \triangleleft R_j = \phi$ if $\forall x$ in $R_i \forall y$ in $R_j : \neg (p_i(x) \land p_j(y))$

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Reduction for PHF

Reduction with join -- Example

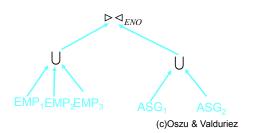
- Assume EMP fragmented as before, and

$$ASG_1 = \sigma_{ENO \leq "E3"}(ASG)$$

$$ASG_2 = \sigma_{ENO>"E3"}(ASG)$$



 SELECT * FROM EMP,ASG WHERE EMP.ENO=ASG.ENO

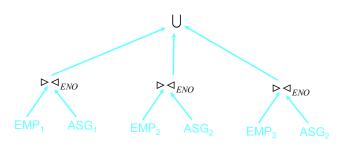


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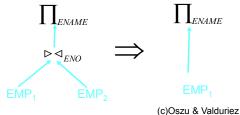
Reduction for PHF

- Reduction with join -- Example
 - Distribute join over unions
 - Apply the reduction rule



Reduction for VF

- Find useless (not empty) intermediate relations
 - Relation R defined over attributes A={A1, ..., An} vertically fragmented as *Ri*= Π_{A'}(*R*) where A' is a subset of *A*
 - $-P_{D,K}(R_i)$ is useless if D is not in A'
 - Example $EMP_1 = \Pi_{ENO, ENAME}(EMP)$, $EMP_2 = \Pi_{ENO, TITLE}(EMP)$
 - SELECT ENAME FROM EMP



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Reduction for DHF

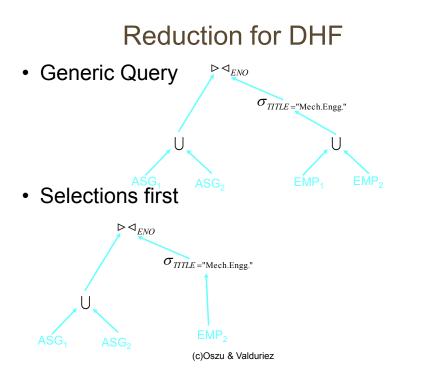
- Rule:
 - Distribute join over unions
 - Apply the join reduction for horizontal fragmentation
 - Example

$$ASG_1 = ASG \triangleright <_{ENO} (EMP_1)$$

$$ASG_2 = ASG \triangleright <_{ENO} (EMP_2)$$

$$EMP_1 = \sigma_{TTTLE = "Programmetr}(EMP)$$

$$EMP_2 = \sigma_{TTTLE \neq "Programmetr}(EMP)$$
SELECT
$$*$$
FROM
$$EMP, ASG$$
WHERE
$$ASG.ENO=EMP.ENO$$
AND
$$EMP.TITLE="Mech. Engg"$$



Step 3 – Global Optimization

- Input: Fragment query
- Find the best (not necessarily optimal) global schedule
 - Minimize a cost function
 - Distributed join processing
 - Bushy vs. linear trees
 - Which relation to ship where?
 - · Ship-whole vs. ship-as-needed
 - Decide on use of semijoins
 - Join methods
 - Nested loop vs. ordered joins (merge join or hash join)

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Semijoin Algorithms

- Perform the join
 - Send R to site 2
 - Site 2 computes the join
- Consider semijoin $R \triangleright \triangleleft_A S \Leftrightarrow (R \triangleright \triangleleft_A S) \triangleright \triangleleft_A S$
 - $\neg S' \leftarrow \prod_A(S)$
 - $-S' \rightarrow Site 1$
 - Site 1 computes $R' = R \triangleright <_A S'$
 - $-R' \rightarrow$ Site 2

- Site 2 computes $R' \triangleright \triangleleft_A S$

Semijoin is better if

 $size(\prod_{A}(S)) + size(R \triangleright <_{A} S)) < size(R)$

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R* Algorithm

- Performing Joins
- Ship Whole
 - Larger data transfer
 - Smaller number of messages
 - Better if relation are small
- Fetch as needed
 - Number of message O(card of external relation)
 - Data transfer per message is minimal
 - Better if relations are large and selectivity is good.