

Topic 3: Efficient query evaluation

Unit 1: Acyclic query evaluation

Lecture 19

Wolfgang Gatterbauer

CS7240 Principles of scalable data management (sp24)

<https://northeastern-datalab.github.io/cs7240/sp24/>

3/26/2024

Pre-class conversations

- Last class summary
- Project: we can iterate, just approach me (P3 today: TUE, 3/26)
- Scribes: we are past halftime of the class (0-5 / 7)
 - please see my detailed comments, do also approach me after class with comments / questions / pointers
 - I will have caught up by next week TUE
- Today:
 - the semi-join reduction as basis for efficient algorithms

Topic 3: Efficient Query Evaluation & Factorized Representations

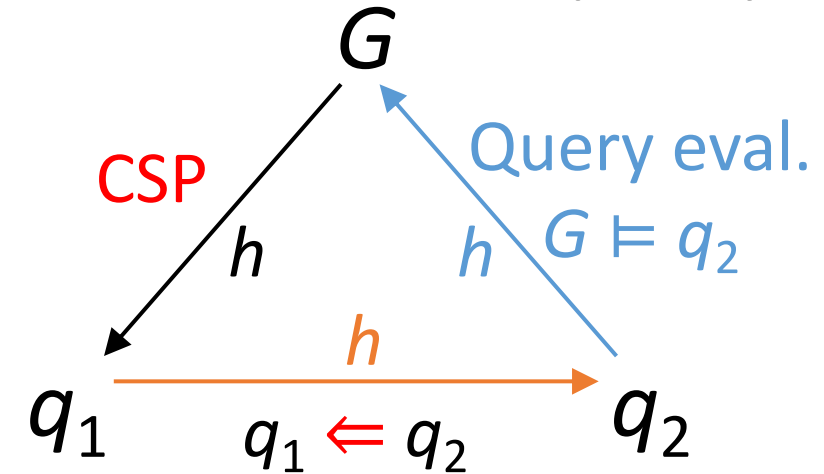
- ~~Lecture 19 (Fri 3/22): T3-U1 Acyclic Queries~~
- **Lecture 20 (Tue 3/26): T3-U2 Cyclic Queries**
- **Lecture 21 (Fri 3/29): T3-U2 Cyclic Queries**
- **Lecture 22 (Tue 4/2): T3-U2 Cyclic Queries**
- **Lecture 23 (Fri 4/5): T3-U3 Factorized Representations**
- **Lecture 24 (Tue 4/9): T3-U4 Optimization Problems & Top-k**

Pointers to relevant concepts & supplementary material:

- **Unit 1. Acyclic Queries:** query hypergraph, Yannakakis algorithm, GYO reduction, dynamic programming, algebraic semirings, [Alice] Ch6.4, [Koutris'19] L4, enumeration, ranked enumeration:[Tziavelis+'20]
- **Unit 2. Cyclic Queries:** tree & hypertree decomposition, query widths, fractional hypertree width, AGM bound, worst-case optimal join algorithms, optimal algorithms, submodular width and multiple decompositions: [AGM'13], [NPRR'18], [KNR'17], [KNS'17], [Gottlob+'16]
- **Unit 3. Factorized Representations:** normalization, factorized databases [Olteanu, Schleich'16]
- **Unit 4. Optimization Problems & Top-k:** shortest paths, dynamic programming (DP), Yannakakis, semirings, rankings, top-k: [Roughgarden'10], [Ilyas+08], [Rahul, Tao'19], ranked enumeration [Tziavelis+'20a], [Tziavelis+'20b]

Islands of Tractability of CQ Evaluation

- Major Research Program: Identify tractable cases of the combined complexity of conjunctive query evaluation.
- Over the years, this program has been pursued by two different research communities:
 - The **Database Theory community**
 - The **Constraint Satisfaction community**
- Explanation: Problems in those community are closely related:



Constraint Satisfaction Problem \equiv Homomorphism Problem \equiv CQ evaluation

[Feder, Vardi 1993]

[Chandra, Merlin 1977]

[Kolaitis, Vardi 2000]

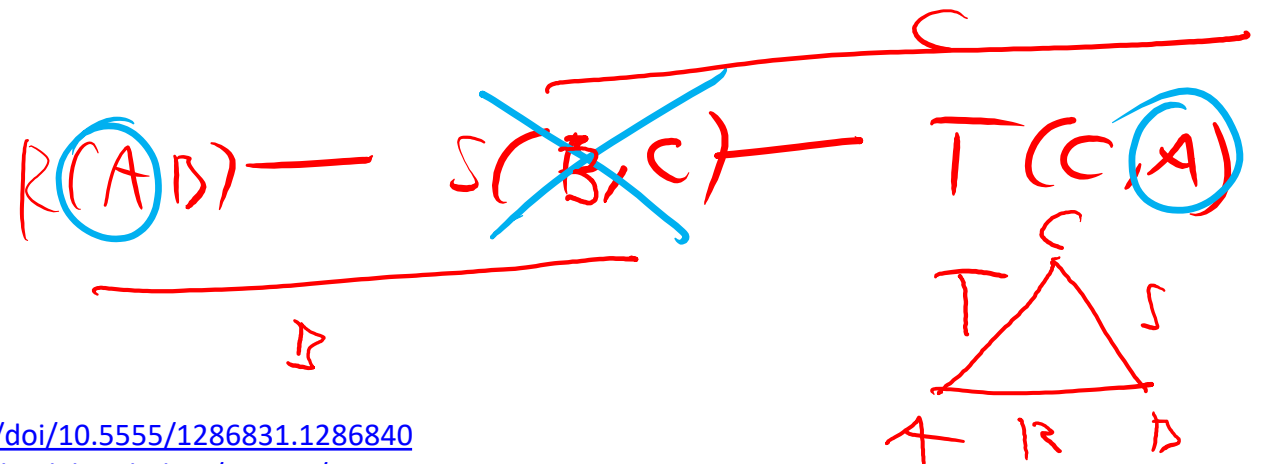
Feder, Vardi: Monotone monadic SNP and constraint satisfaction, STOC 1993 <https://doi.org/10.1145/167088.167245> / Kolaitis, Vardi: Conjunctive-Query Containment and Constraint Satisfaction, JCSS 2000 <https://doi.org/10.1006/jcss.2000.1713> / Chandra, Merlin. "Optimal implementation of conjunctive queries in relational data bases", STOC 1977. <https://doi.org/10.1145/800105.803397>

Based on Phokion Kolaitis' "Logic and Databases" series at Simons Institute, 2016. <https://simons.berkeley.edu/talks/logic-and-databases>

Wolfgang Gatterbauer. Principles of scalable data management: <https://northeastern-datalab.github.io/cs7240/>

Acyclic queries

- Like many areas in computer science, cycles complicate things
 - Same with conjunctive query (CQ) evaluation
- **Acyclic CQs** are a large and useful tractable case CQs
- A query is **acyclic** if its relations can be placed in a tree (**join tree**) s.t.
 - the set of nodes that contain any variable form a connected set
- **Yannakakis' algorithm** [81]: any acyclic query can be computed in time: $O(|\text{Input}| + |\text{Output}|)$



Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

<https://northeastern-datalab.github.io/>

Semijoin (\bowtie): derived RA operator



Actor (aid, name, gender)
Play (aid, mid, role)
Movie(mid, name, year)

- $R \bowtie S$: Find tuples in R for which there is a matching tuple in S that is equal on their common attribute names.

$$R \bowtie S = \pi_{A_1, \dots, A_n} (R \Join S) \quad \text{RA}$$

where A_1, \dots, A_n are the attributes in R

Intuition: remove "dangling tuples" in R

- Example:
"Find actors who play some role."

RA: ?

Semijoin (\bowtie): derived RA operator



Actor (aid, name, gender)
Play (aid, mid, role)
Movie(mid, name, year)

- $R \bowtie S$: Find tuples in R for which there is a matching tuple in S that is equal on their common attribute names.

$$R \bowtie S = \pi_{A_1, \dots, A_n} (R \Join S) \quad \text{RA}$$

where A_1, \dots, A_n are the attributes in R

SQL: ?

Intuition: remove "dangling tuples" in R

- Example:
"Find actors who play some role."

RA: Actor \bowtie Play

Semijoin (\bowtie): derived RA operator



Actor (aid, name, gender)
Play (aid, mid, role)
Movie(mid, name, year)

- $R \bowtie S$: Find tuples in R for which there is a matching tuple in S that is equal on their common attribute names.

$$R \bowtie S = \pi_{A_1, \dots, A_n} (R \Join S) \quad \text{RA}$$

where A_1, \dots, A_n are the attributes in R

Intuition: remove "dangling tuples" in R

- Example:
"Find actors who play some role."

RA: Actor \bowtie Play

Semijoins have no "direct" representation in SQL
SQL: (just like relational division)

```
SELECT DISTINCT *  
FROM Actor  
WHERE aid IN  
      (SELECT aid FROM Play)
```

Alternative:
WHERE EXISTS

```
SELECT DISTINCT *  
FROM Actor A  
LEFT JOIN Play P  
ON A.aid = P.aid  
WHERE P.aid IS NOT NULL
```

```
SELECT DISTINCT  
      A.aid, name, age  
FROM Actor A, Play P  
WHERE A.aid = P.aid
```

Are those
3 queries
equivalent



Semijoin (\bowtie): derived RA operator



Actor (aid, name, gender)
Play (aid, mid, role)
Movie(mid, name, year)

- $R \bowtie S$: Find tuples in R for which there is a matching tuple in S that is equal on their common attribute names.

$$R \bowtie S = \pi_{A_1, \dots, A_n} (R \Join S) \quad \text{RA}$$

where A_1, \dots, A_n are the attributes in R

Intuition: remove "dangling tuples" in R

- Example:
"Find actors who play some role."

RA: Actor \bowtie Play

Semijoins have no "direct" representation in SQL
SQL: (just like relational division)

```
SELECT DISTINCT *  
FROM Actor  
WHERE aid IN  
      (SELECT aid FROM Play)
```

```
SELECT DISTINCT *  
FROM Actor A  
LEFT JOIN Play P  
ON A.aid = P.aid  
WHERE P.aid IS NOT NULL
```

```
SELECT DISTINCT  
      A.aid, name, age  
FROM Actor A, Play P  
WHERE A.aid = P.aid
```

Duplicates in R are preserved!

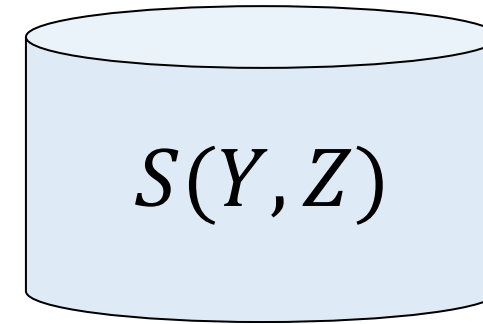
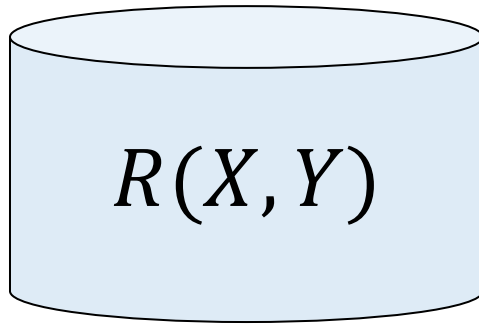
Equivalent only if no duplicates in Actor!

Semijoins in Distributed Databases

- Semijoins are often used to compute equijoins in distributed databases

$R \bowtie S$

Goal: send less data to reduce network bandwidth!

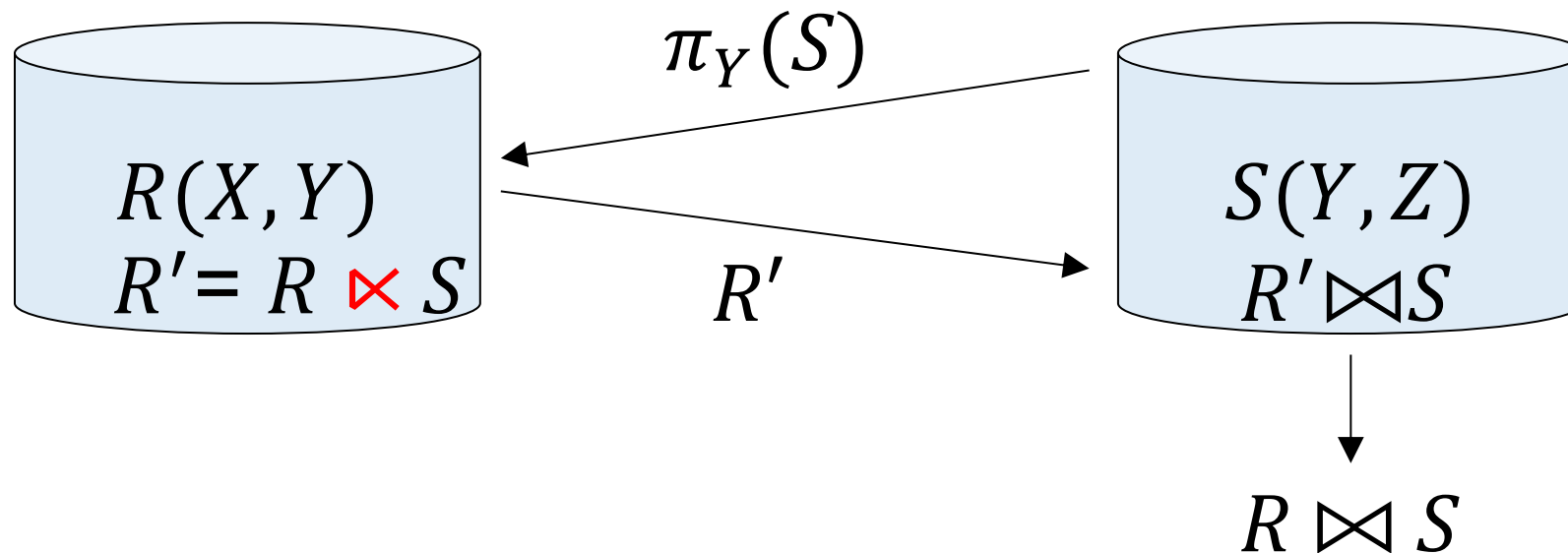


Semijoins in Distributed Databases

- Semijoins are often used to compute equijoins in distributed databases

$R \bowtie S$

Goal: send less data to reduce network bandwidth!



$R \bowtie S = (R \ltimes S) \bowtie S$ law of semijoins

reduced

Semijoins in Distributed Databases

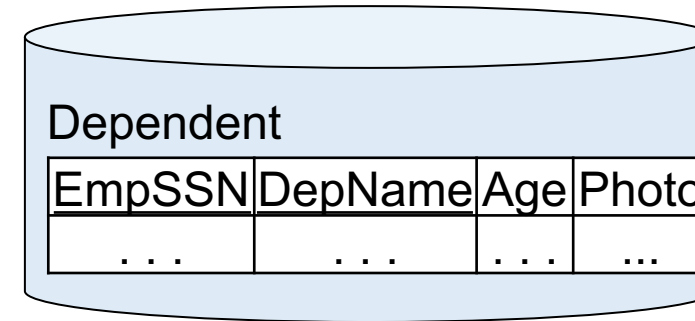
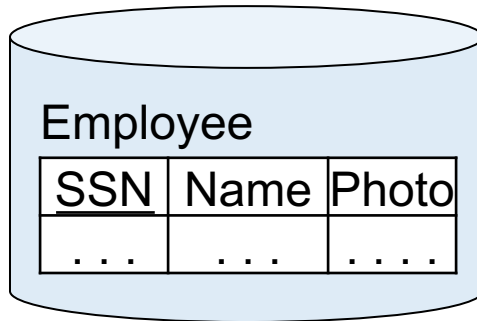


Task: compute with minimum data transfer:

Employee $\bowtie_{\text{SSN}=\text{EmpSSN}} (\sigma_{\text{age}>71}(\text{Dependent}))$

Assumptions:

1. Very few employees have dependents.
2. Very few dependents have age > 71.
3. "Photo" is big.



?

Semijoins in Distributed Databases

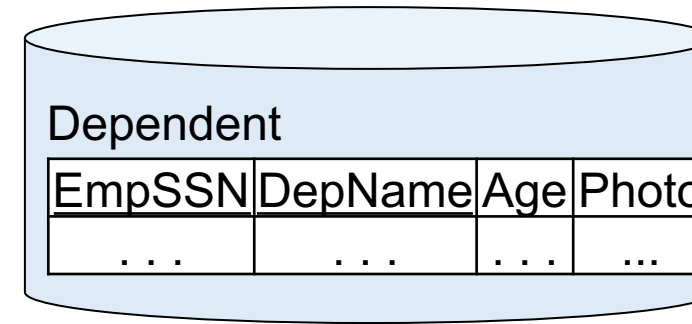
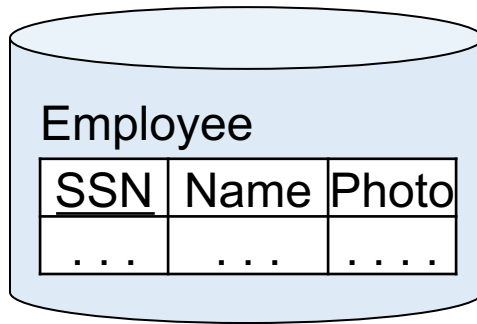


Task: compute with minimum data transfer:

$\text{Employee} \bowtie_{\text{SSN}=\text{EmpSSN}} (\sigma_{\text{age}>71}(\text{Dependent}))$

Assumptions:

1. Very few employees have dependents.
2. Very few dependents have age > 71.
3. "Photo" is big.



R

$R(\text{SSN}) = \pi_{\text{EmpSSN}} \sigma_{\text{age}>71}(\text{Dependents})$

"R": message from the right side

$L = \text{Employee} \bowtie_{\text{SSN}=\text{EmpSSN}} R$
 $= \text{Employee} \bowtie_{\text{SSN}=\text{EmpSSN}} (\sigma_{\text{age}>71}(\text{Dependent}))$
 $= \text{Employee} \bowtie_{\text{SSN}=\text{EmpSSN}} R$

"L": message from the leftside

L

$\text{Answer} = L \bowtie_{\text{SSN}=\text{EmpSSN}} \text{Dependent}$

Semijoins in Distributed Databases

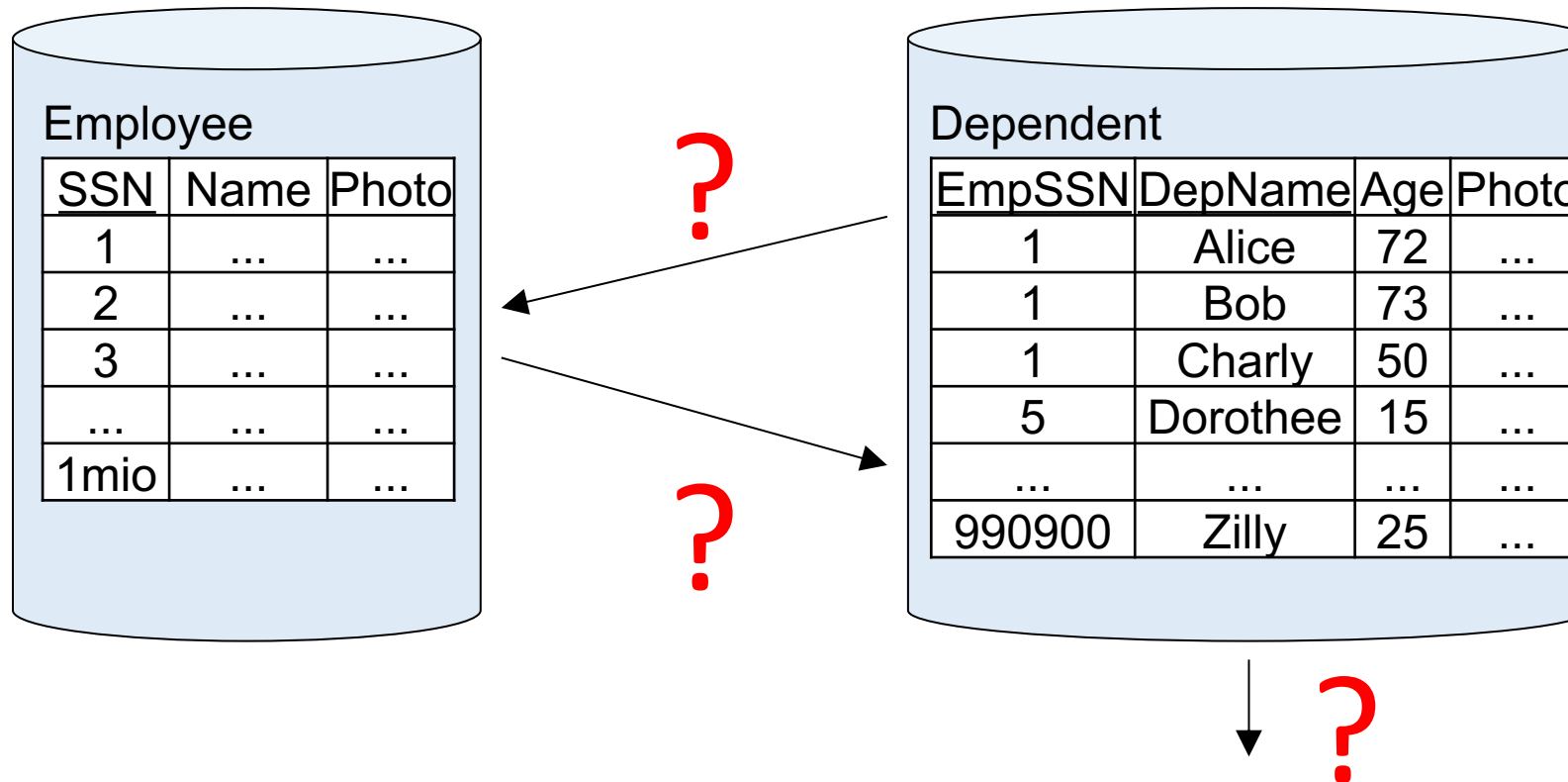


Task: compute with minimum data transfer:

Employee $\bowtie_{SSN=EmpSSN} (\sigma_{age>71}(\text{Dependent}))$

Assumptions:

1. Very few employees have dependents.
2. Very few dependents have age > 71.
3. "Photo" is big.



Semijoins in Distributed Databases

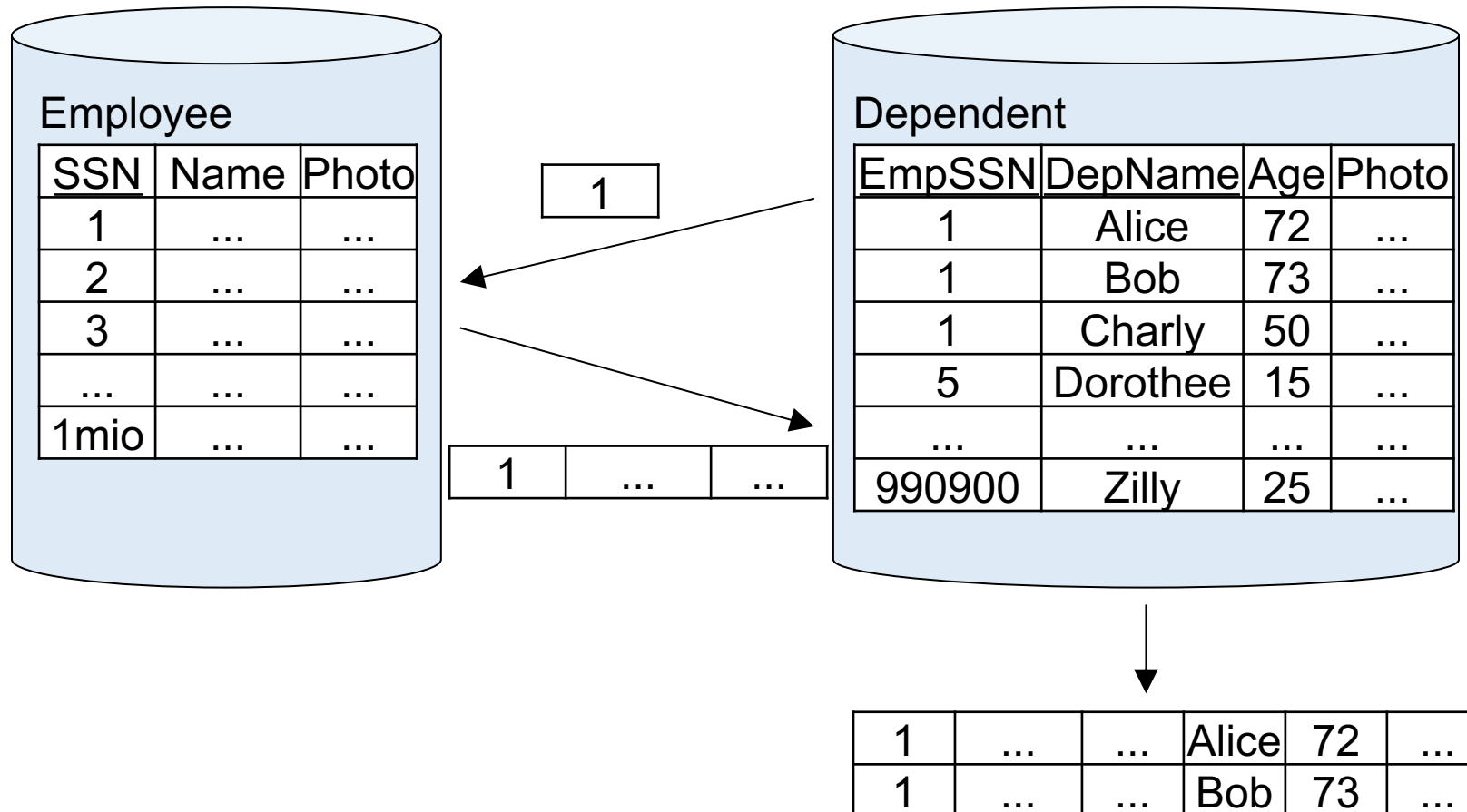


Task: compute with minimum data transfer:

$\text{Employee} \bowtie_{\text{SSN}=\text{EmpSSN}} (\sigma_{\text{age}>71}(\text{Dependent}))$

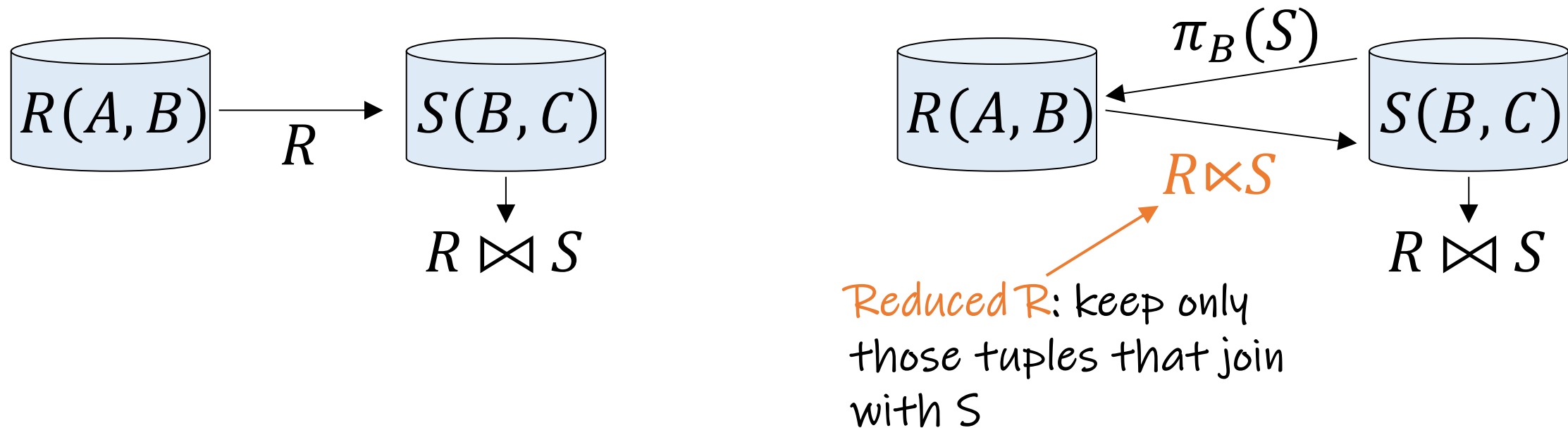
Assumptions:

1. Very few employees have dependents.
2. Very few dependents have age > 71.
3. "Photo" is big.



Semijoins as Message Passing

- **Semijoins** can reduce network use for equijoins in distributed databases



Effective if 1) the size of join attribute B (or number of distinct values) is smaller than A and C , and 2) few tuples from R participate in the join

Repetition: Law of Semijoins

- Definition: the **semi-join** operation is:

$$R \bowtie S = \pi_{A_1, \dots, A_n}(R \bowtie S)$$

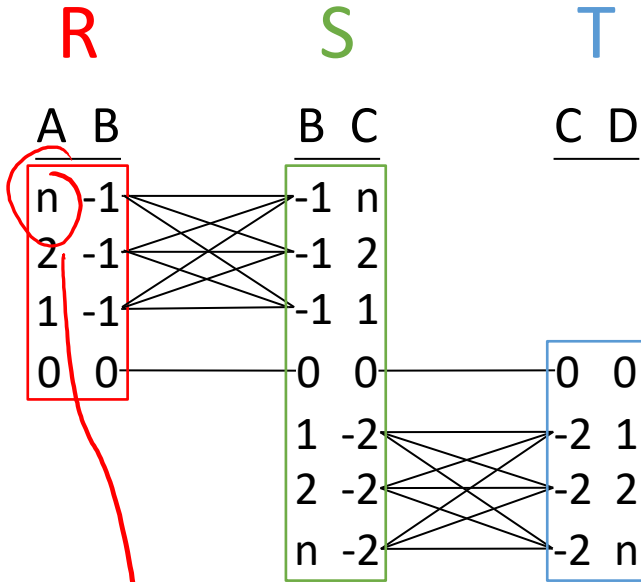
- Formally, $R \bowtie S$ means: retain from R only those tuples that have some matching tuple in S (in bag semantics: duplicates in R are preserved / Duplicates in S don't matter)
 - Data complexity: $O(|R| + |S|)$ ignoring log-factors
 - Input: $R(A_1, \dots, A_n)$, $S(B_1, \dots, B_m)$, Output: $T(A_1, \dots, A_n)$
- The **law of semijoins** is:

$$R \bowtie S = (R \bowtie S) \bowtie S$$

- Thus, **removing "dangling tuples"** from a table does not change the query result

Semi-joins can also help if data are local

$$Q_3^\infty(x, y, z, x) : -R(x, y), S(y, z), T(z, w)$$



$$Q = (R \bowtie_B S) \bowtie_C T$$

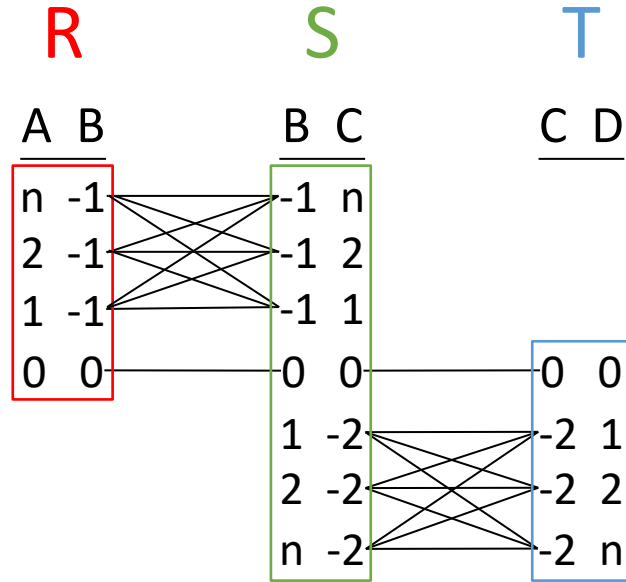
What is the problem?

?

3, 5, 100, 1000, 104, ..

Semi-joins can also help if data are local

$$Q_3^\infty(x, y, z, x) : -R(x, y), S(y, z), T(z, w)$$



$$Q = R \bowtie_B S \bowtie_C T$$

What is the problem?

- Query output cardinality is 1
- But quadratic intermediate result sizes

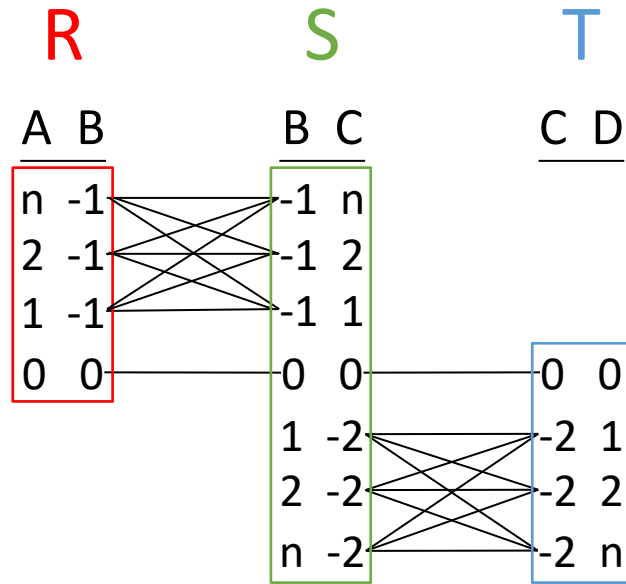
Thus the query takes $O(n^2)$ despite constant output ☹

What is a typical query plan?



Semi-joins can also help if data are local

$$Q_3^\infty(x, y, z, x) : -R(x, y), S(y, z), T(z, w)$$



$$Q = R \bowtie_B S \bowtie_C T$$

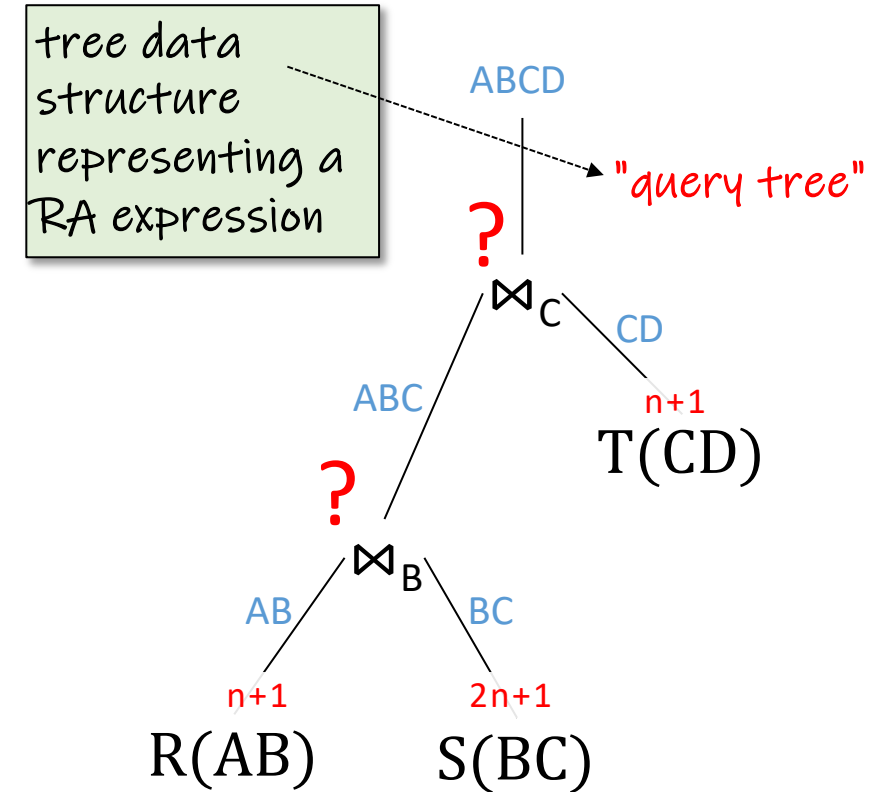
What is the problem?

- Query output cardinality is 1
- But quadratic intermediate result sizes

Thus the query takes $O(n^2)$ despite constant output ☹

What is a typical query plan?

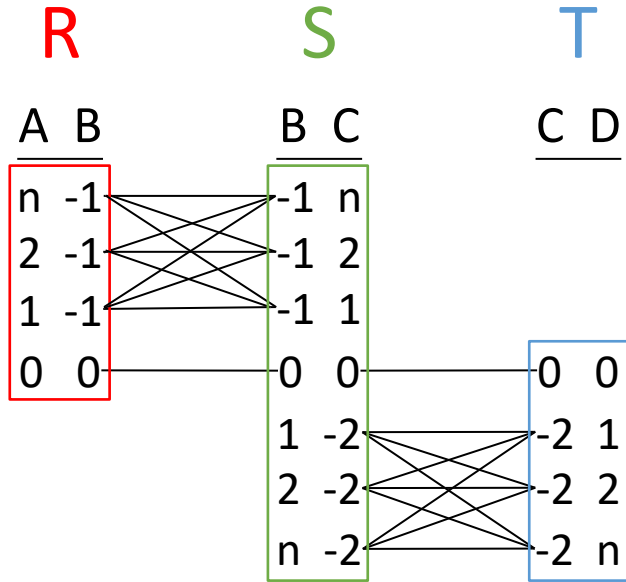
$$Q = ((R \bowtie_B S) \bowtie_C T)$$



What are the cardinalities at each stage?

Semi-joins can also help if data are local

$$Q_3^\infty(x, y, z, x) : -R(x, y), S(y, z), T(z, w)$$



$$Q = R \bowtie_B S \bowtie_C T$$

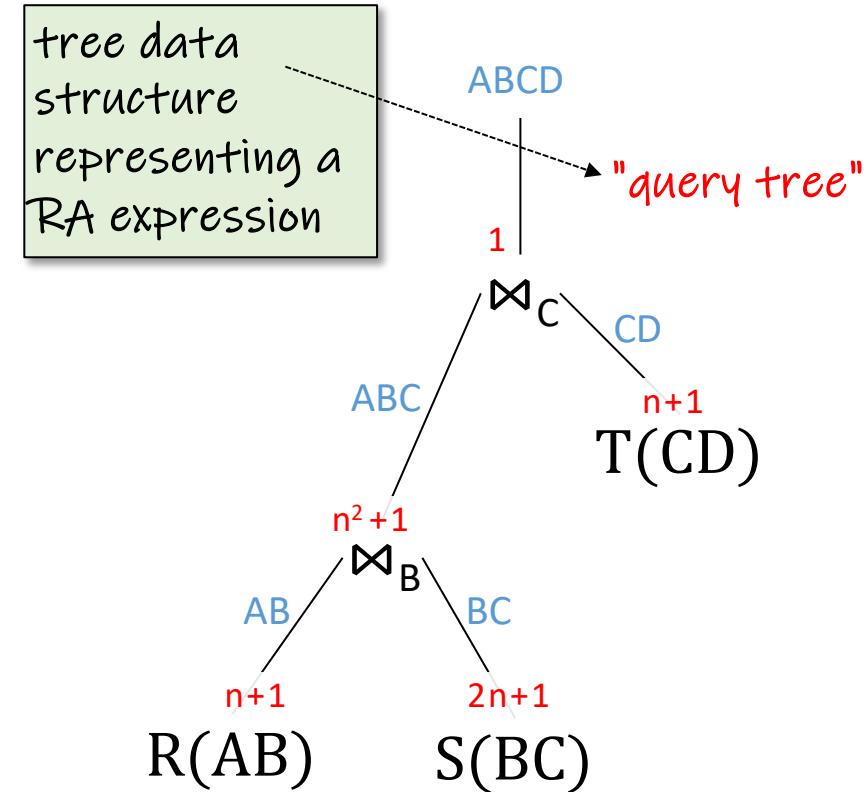
What is the problem?

- Query output cardinality is 1
- But quadratic intermediate result sizes

Thus the query takes $O(n^2)$ despite constant output ☹

What is a typical query plan?

$$Q = ((R \bowtie_B S) \bowtie_C T)$$

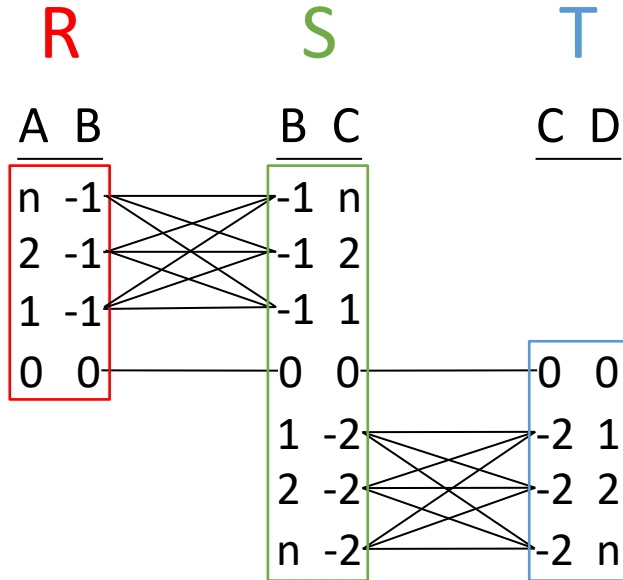


What are the cardinalities at each stage?

Semi-joins can also help if data are local



$Q_3^\infty(x, y, z, x): -R(x, y), S(y, z), T(z, w)$



?

Can you think of a faster evaluation plan?

-- Query 1

```
select *  
into record1  
from R natural join S natural join T;
```

$n=1,000: t_{Q1}=1.4 \text{ sec}$

$n=2,000: t_{Q1}=6.1 \text{ sec}$

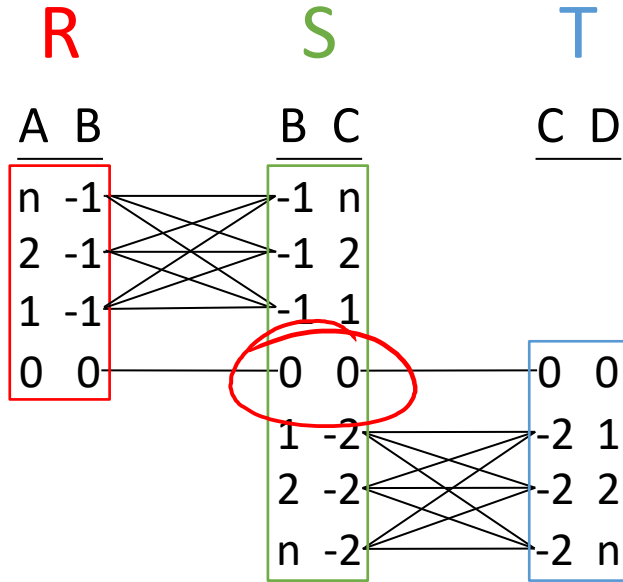
$t_{Q2}=5 \text{ msec}$

$t_{Q2}=8 \text{ msec}$

Semi-joins can also help if data are local



$Q_3^\infty(x, y, z, x): -R(x, y), S(y, z), T(z, w)$



WITH clauses (also CTE = Common Table Expression) act like temporary views

Only 1 single output tuple!

-- Query 1

```
select *
into record1
from R natural join S natural join T;
```

$n=1,000: t_{Q1}=1.4 \text{ sec}$

$n=2,000: t_{Q1}=6.1 \text{ sec } O(n^2) \text{ ☹️}$

-- Query 2

```
With S2 as
  (SELECT *
   FROM S
   WHERE S.B in
     (SELECT R.B
      FROM R)),
S3 as
  (SELECT *
   FROM S2
   WHERE S2.C in
     (SELECT T.C
      FROM T))
select a, b, c, d
into record2
from R natural join S3 natural join T;
```

$t_{Q2}=5 \text{ msec}$

$t_{Q2}=8 \text{ msec } O(n) \text{ ☺️}$

Diversion into CTE's (Common Table Expressions)



```
select *
into record3
from R natural join
```

```
(SELECT *
FROM
  (SELECT *
   FROM S
   WHERE S.B in
     (SELECT R.B
      FROM R)) S2
WHERE S2.C in
  (SELECT T.C
   FROM T)) S3
```

```
natural join T;
```

WITH clauses (also **CTE** = Common Table Expression) act like temporary views

- Allow to deconstruct more complex queries into simple blocks to be used and reused if necessary.
- Can increase readability by emphasizing a more procedural interpretation of a query in a workflow
- Especially useful if you need to reference a derived table multiple times in a single query or you perform the same calculation multiple times across multiple query components (= memoization)

-- Query 2

With S2 as

```
(SELECT *
FROM S
WHERE S.B in
  (SELECT R.B
   FROM R)),
```

S3 as

```
(SELECT *
FROM S2
WHERE S2.C in
  (SELECT T.C
   FROM T))
```

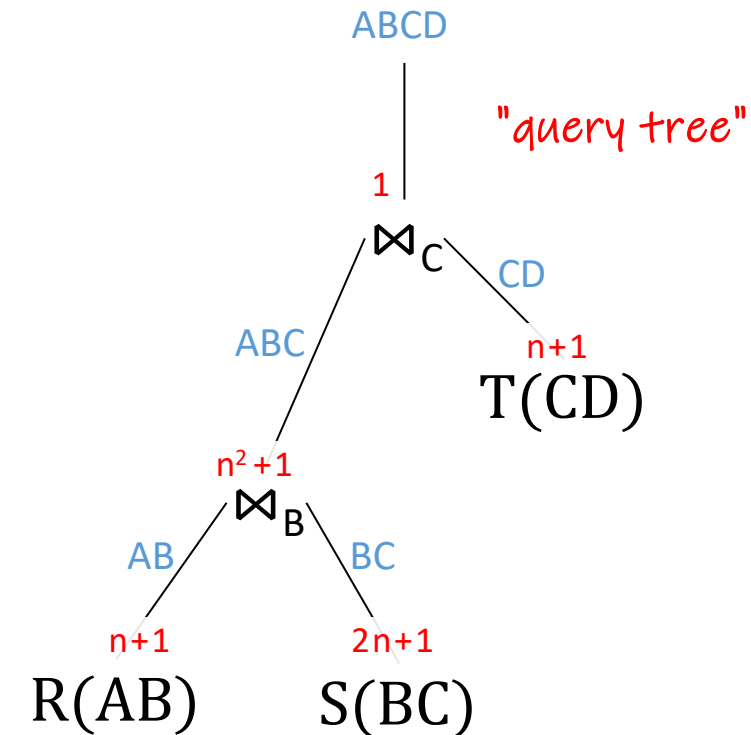
```
select a, b, c, d
into record2
from R natural join S3 natural join T;
```


The more general idea: "Sideways information passing"

Sideways information passing:

- "sending information from one subexpression not simply to its parent expression, but also to some other correlated portion of the query computation, in order to prune irrelevant results" [Ives, Taylor 08]
- includes techniques like **two-way semijoins** [Bernstein, Goodman 81] and **magic sets** [Beeri, Ramakrishnan 91]

$$Q = ((R \bowtie_B S) \bowtie_C T)$$



[Bernstein, Goodman 81]. "Using Semi-Joins to Solve Relational Queries", JACM 1981. <https://doi.org/10.1145/322234.322238>

[Beeri, Ramakrishnan 91]: "On the power of magic", Journal of Logic Programming, 1991. [https://doi.org/10.1016/0743-1066\(91\)90038-Q](https://doi.org/10.1016/0743-1066(91)90038-Q)

Definition from: [Ives, Taylor 08]. "Sideways Information Passing for Push-Style Query Processing", ICDE 2008. <https://doi.org/10.1109/ICDE.2008.4497486>

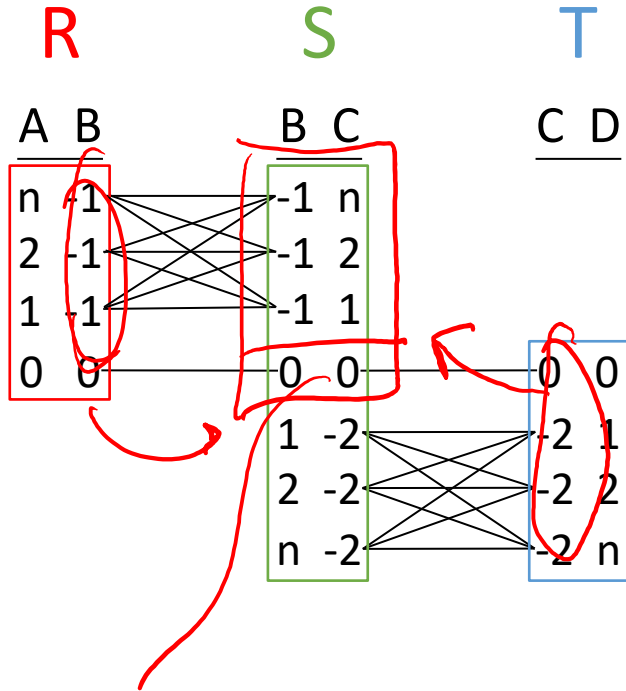
Wolfgang Gatterbauer. Principles of scalable data management: <https://northeastern-datalab.github.io/cs7240/>

The more general idea: "Sideways information passing"

$Q_3^\infty(x, y, z, x): -R(x, y), S(y, z), T(z, w)$



601



-- Query 2

With S2 as

```
(SELECT *
FROM S
WHERE S.B in
  (SELECT R.B
   FROM R)),
```

S3 as

```
(SELECT *
FROM S2
WHERE S2.C in
  (SELECT T.C
   FROM T))
```

```
select a, b, c, d
into record2
from R natural join S3 natural join T;
```

$\{0, -1\}$
 $S(0, 0)$

Topic 3: Efficient query evaluation

Unit 1: Acyclic query evaluation

Lecture 20

Wolfgang Gatterbauer

CS7240 Principles of scalable data management (sp24)

<https://northeastern-datalab.github.io/cs7240/sp24/>

3/29/2024

Pre-class conversations

- Last class summary
- Project: Feedback by next week TUE
 - we can iterate, just approach me
- Scribes: Feedback by next week TUE (end of day)
 - please see my detailed comments, do also approach me after class with comments / questions / pointers. The goal is for the scribes and comments to be starters

- Today:
 - from semi-join to Yannakakis, via (alpha-)acyclic queries

Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

<https://northeastern-datalab.github.io/>

Outline journey

- The idea from the previous slides on **semijoins** can be generalized to more than two joins, as long as the joins are "**acyclic**"
- We will next define "**acyclic queries**"
 - It is a bit more complicated for general hypergraphs (with relations of arbitrary arity) than for graphs
- And later see the "**semi-join reduction**" on general acyclic queries
- Which is basically the Yannakakis algorithm

Query Hypergraph (vs. Dual Hypergraph vs. Incidence Matrix)

$Q: -A(x), R(x, v), S(v, y), B(y, u), T(y, z), C(z), D(z, w)$

Query hypergraph (nodes = variables)

?

Incidence matrix

?

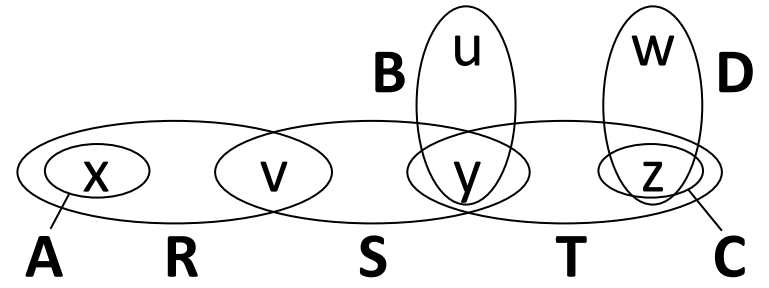
Query dual hypergraph (nodes = atoms)

?

Query Hypergraph (vs. Dual Hypergraph vs. Incidence Matrix)

$Q: \neg A(x), R(x, v), S(v, y), B(y, u), T(y, z), C(z), D(z, w)$

Query hypergraph (nodes = variables)



determines α -acyclicity of CQs

Incidence matrix



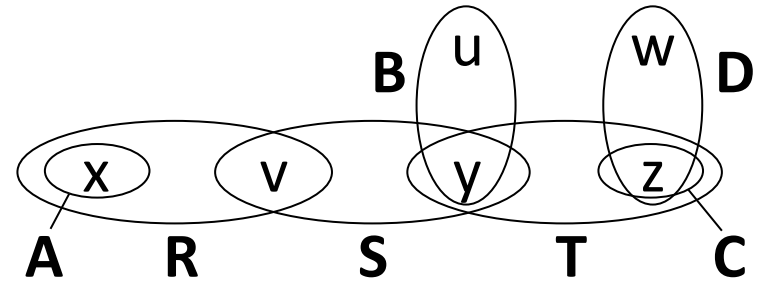
Query dual hypergraph (nodes = atoms)



Query Hypergraph (vs. Dual Hypergraph vs. Incidence Matrix)

$Q: -A(x), R(x, v), S(v, y), B(y, u), T(y, z), C(z), D(z, w)$

Query hypergraph (nodes = variables)

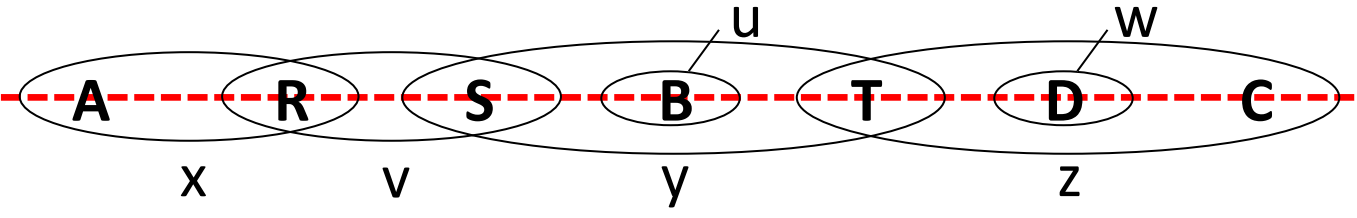


determines α -acyclicity of CQs

Incidence matrix



Query dual hypergraph (nodes = atoms)

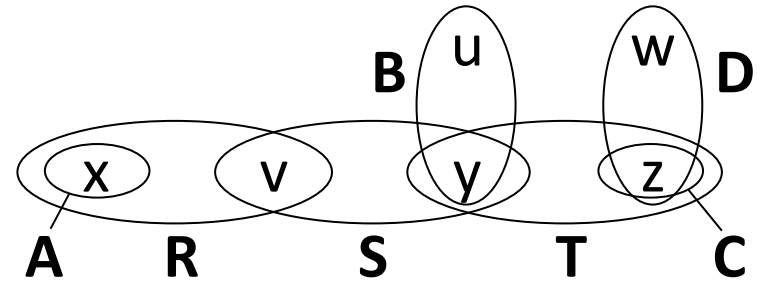


determines complexity of resilience [Meliou+'10]

Query Hypergraph (vs. Dual Hypergraph vs. Incidence Matrix)

$Q: -A(x), R(x, v), S(v, y), B(y, u), T(y, z), C(z), D(z, w)$

Query hypergraph (nodes = variables)

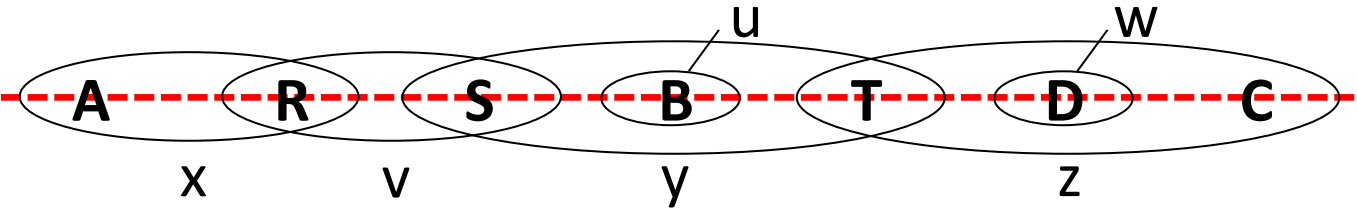


determines α -acyclicity of CQs

Incidence matrix

determines minimal query plans [G+'17] (intuitively, plans with early projections)

Query dual hypergraph (nodes = atoms)



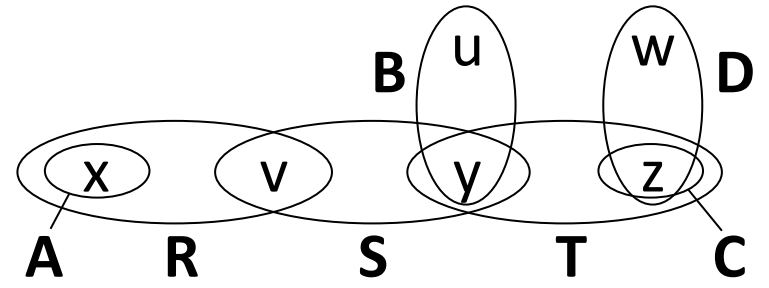
determines complexity of resilience [Meliou+'10]

	x	v	y	u	z	w
A	1					
R	1	1				
S		1	1			
B			1	1		
T			1		1	
C					1	
D					1	1

Query Hypergraph (vs. Dual Hypergraph vs. Incidence Matrix)

$Q: -A(x), R(x, v), S(v, y), B(y, u), T(y, z), C(z), D(z, w)$

Query hypergraph (nodes = variables)



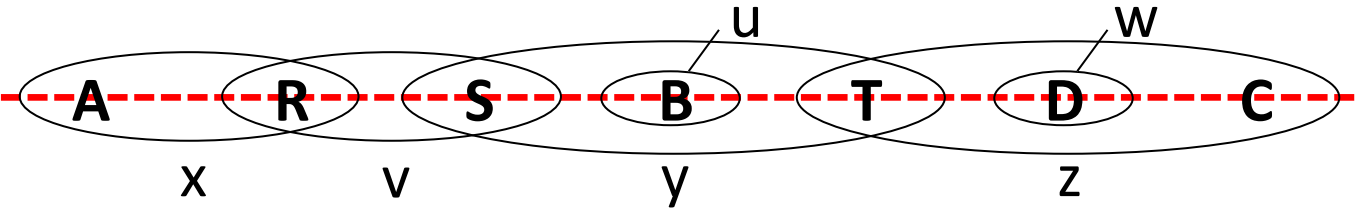
determines α -acyclicity of CQs

Incidence matrix

determines minimal query plans [G+'17] (intuitively, plans with early projections)

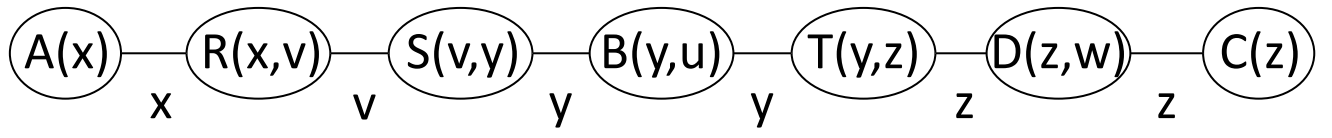
	x	v	y	u	z	w
A	1					
R	1	1				
S		1	1			
B			1	1		
T			1		1	
C					1	
D					1	1

Query dual hypergraph (nodes = atoms)



determines complexity of resilience [Meliou+'10]

Join tree

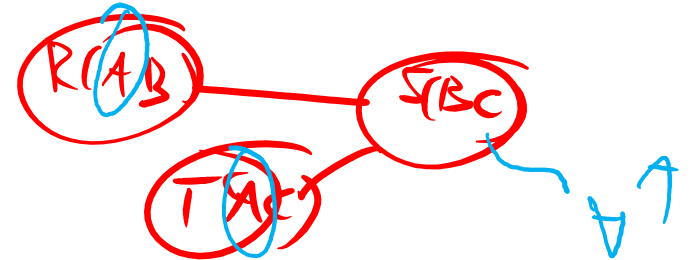


only possible if query is α -acyclic

α (alpha)-acyclic Conjunctive Queries have Join Trees

- Definition: A conjunctive query Q is α -acyclic if it has a **join tree**.
- A **join tree** for $Q(\mathbf{x}) :- R_1(\mathbf{z}_1), R_2(\mathbf{z}_2), \dots, R_m(\mathbf{z}_m)$ is a tree $T = (V, E)$ such that:
 - V : The nodes of T are the **atoms** $R_i(\mathbf{z}_i)$ of Q
 - E : For every variable y occurring in Q , the set of the nodes of T that contain y forms a **(connected) subgraph of T** .
 - Also called **running intersection property** (see also **junction tree algorithm** and **tree decompositions**)
 - in other words, if a variable w occurs in two different atoms $R_j(\mathbf{z}_j)$ and $R_k(\mathbf{z}_k)$ of Q , then it occurs in each atom on the unique path of T joining $R_j(\mathbf{z}_j)$ and $R_k(\mathbf{z}_k)$

$R(A, B), S(B, C), T(A, C)$



- The **GYO reduction*** helps us find a join tree from the hypergraph iff it is α -acyclic

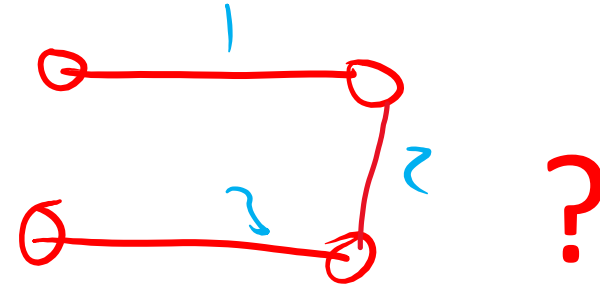
Notice that the definition of join tree does not include the root. We do need to choose roots later, and different choices of roots lead to different sequences of semijoin reductions

* **GYO**: This algorithm is named in honor of Marc H. Graham and the team Clement Yu and Meral Ozsoyoglu, who independently came to essentially this algorithm in [YO79] and [Gra79]: [Gra79] Graham. "On the universal relation." Technical Report, University of Toronto, 1979 / [YO79] Yu, Ozsoyoglu. "An algorithm for tree-query membership of a distributed query." COMPSAC, 1979. <https://doi.org/10.1109/CMPSAC.1979.762509>
Wolfgang Gatterbauer. Principles of scalable data management: <https://northeastern-datalab.github.io/cs7240/>

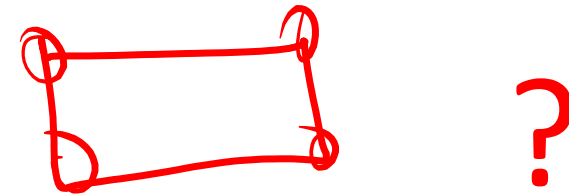
Which query is acyclic?



- Path of length 3? (4 nodes, 3 edges). Return end points.



- Cycle of length 4? Boolean.

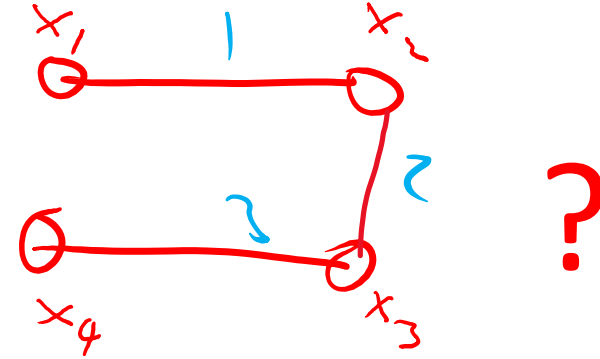


Which query is acyclic?



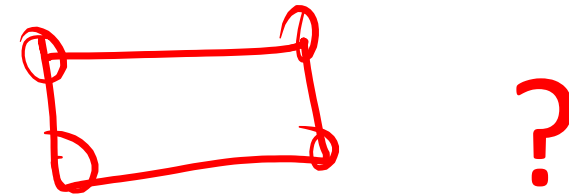
- Path of length 3? (4 nodes, 3 edges). Return end points.

$P4(x_1, x_4) :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4)$



- Cycle of length 4? Boolean.

$C4() :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4), E(x_4, x_1)$

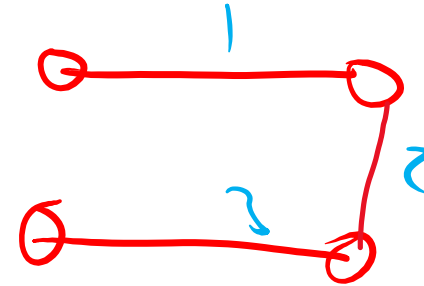


Which query is acyclic?



- Path of length 3? (4 nodes, 3 edges). Return end points.

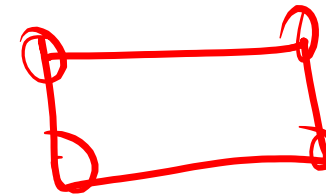
$P4(x_1, x_4) :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4)$



acyclic
(join tree is path)

- Cycle of length 4? Boolean.

$C4() :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4), E(x_4, x_1)$



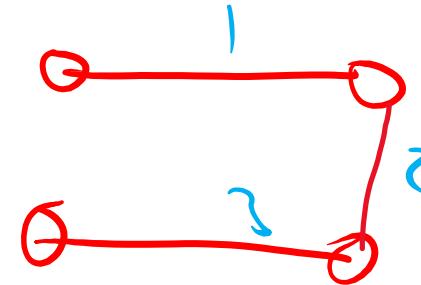
?

Which query is acyclic?



- Path of length 3? (4 nodes, 3 edges). Return end points.

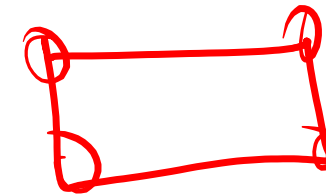
$P4(x_1, x_4) :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4)$



acyclic
(join tree is path)

- Cycle of length 4? Boolean.

$C4() :- E(x_1, x_2), E(x_2, x_3), E(x_3, x_4), E(x_4, x_1)$



cyclic

Acyclic Conjunctive Queries



- Is the following query acyclic?

Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).

Dual Hypergraph (relations as nodes)

Hypergraph (variables as nodes)

?

?

Acyclic Conjunctive Queries

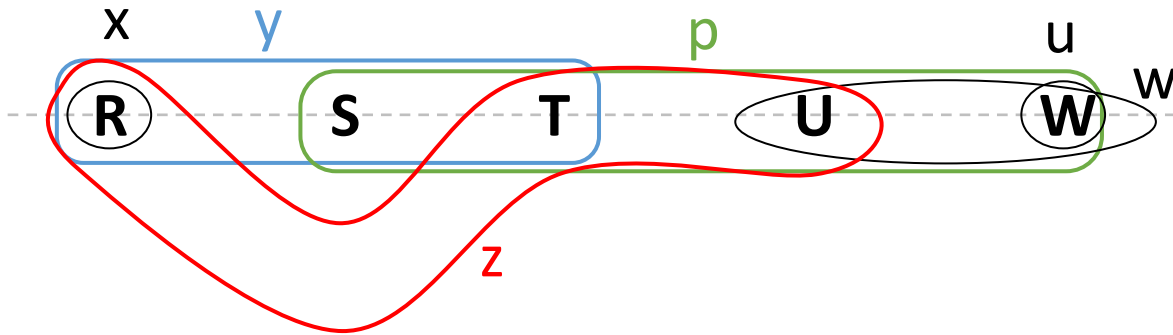


- Is the following query acyclic?

$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$

Dual Hypergraph (relations as nodes)

Hypergraph (variables as nodes)



No linear order in dual hypergraph ☹️
(Thus resilience is NPC!)

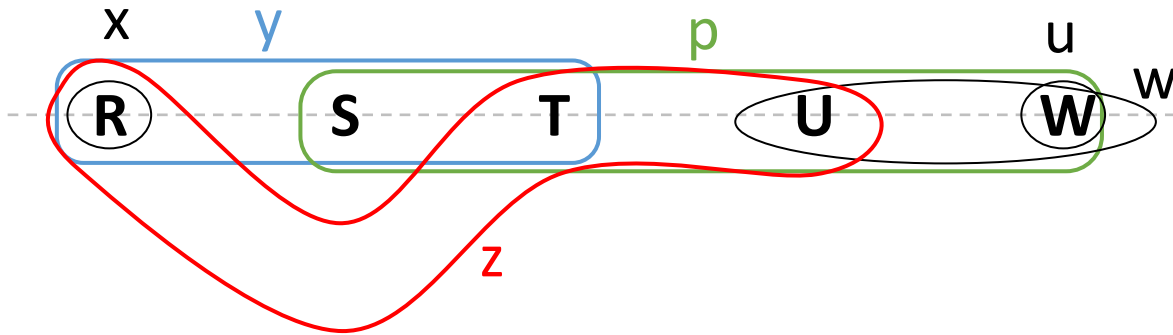
Acyclic Conjunctive Queries



- Is the following query acyclic?

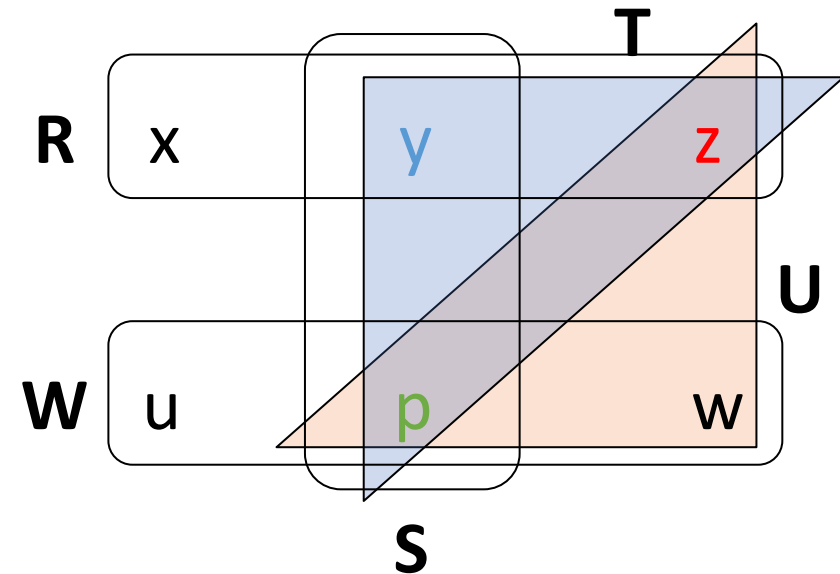
$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$

Dual Hypergraph (relations as nodes)



No linear order in dual hypergraph ☹️
(Thus resilience is NPC!)

Hypergraph (variables as nodes)



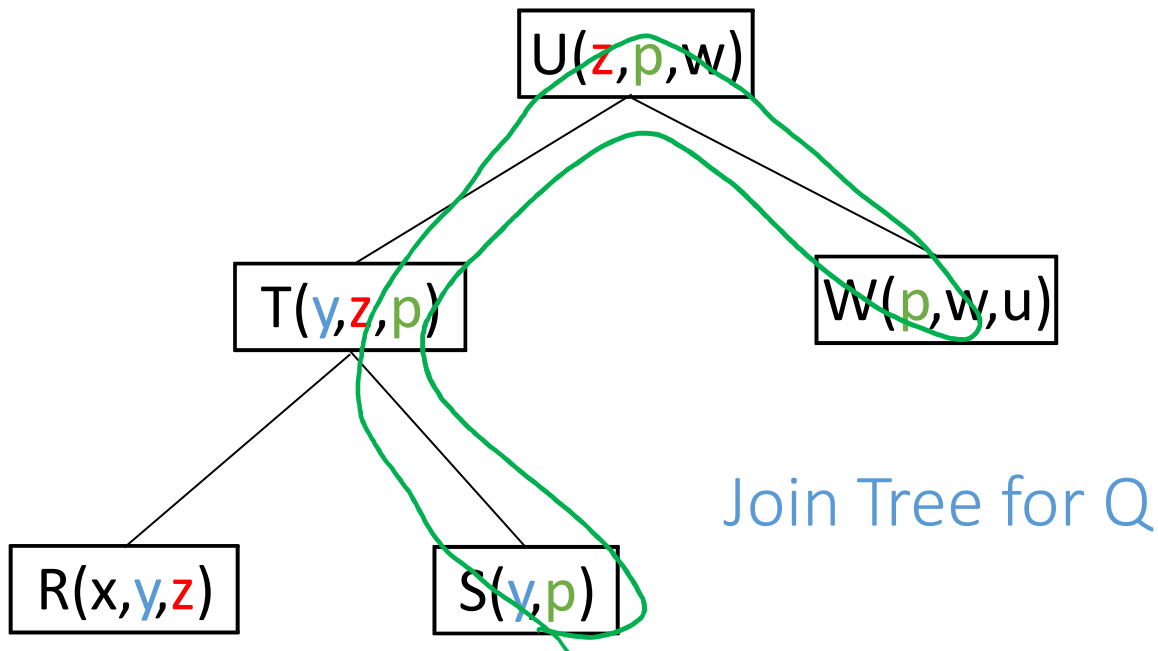
So does the query have a join tree?

Acyclic Conjunctive Queries

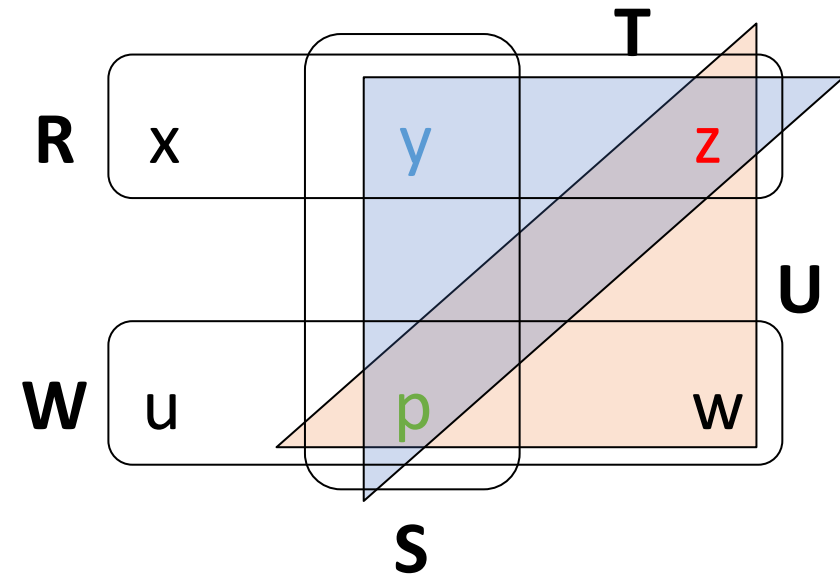


- Is the following query acyclic?

$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$



Hypergraph (variables as nodes)



Yes the query has a join tree

Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

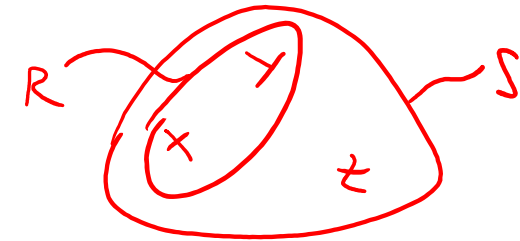
<https://northeastern-datalab.github.io/>

GYO reduction (Graham-Yu-Ozsoyoglu) on the hypergraph

- An **ear** is a hyperedge H whose variables (=nodes) form two groups:
 1. **isolated** variables that appear exclusively in H , and
 2. **join variables** (i.e. they occur in at least one other edge). For join variables there must exist a hyperedge called a **witness** that contains all of them
- **GYO algorithm (Ear removal)**
 - Remove ears greedily from the hypergraph
- Important: any sequence of reductions that removes all hyperedges implies a **join tree**:
 - Just draw an edge between an ear and any witness (notice that if an ear has only isolated nodes, any remaining hyperedge is a witness)

corresponds to a projection

corresponds to a join



This algorithm is named in honor of Marc H. Graham and the team Clement Yu and Meral Ozsoyoglu, who independently came to essentially this algorithm in [YO79] and [Gra79]:
[Gra79] Graham. "On the universal relation." Technical Report, University of Toronto, 1979 / [YO79] Yu, Ozsoyoglu. "An algorithm for tree-query membership of a distributed query." COMPSAC, 1979. <https://doi.org/10.1109/COMPSAC.1979.762509>

Wolfgang Gatterbauer. Principles of scalable data management: <https://northeastern-datalab.github.io/cs7240/>

Proof GYO reduction = acyclic query

- Proof (GYO \Rightarrow acyclic): if GYO leads to an empty hypergraph, then the resulting graph forms a valid join tree
 - notice that by construction, for any variable, those edges containing the variable will form an induced subtree
- Proof (acyclic \Rightarrow GYO): if there is a valid join tree, then removing leaf nodes in any order corresponds to a sequence of GYO reductions:
 - All non-exclusive variables from a leaf must be shared with the parent. Thus the parent forms a witness that **consumes** the leaf (notice that by construction, this also works if the leaf node shares no variables)

GYO reduction: Example 1



$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree

?

Query hypergraph

?

GYO reduction: Example 1



$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

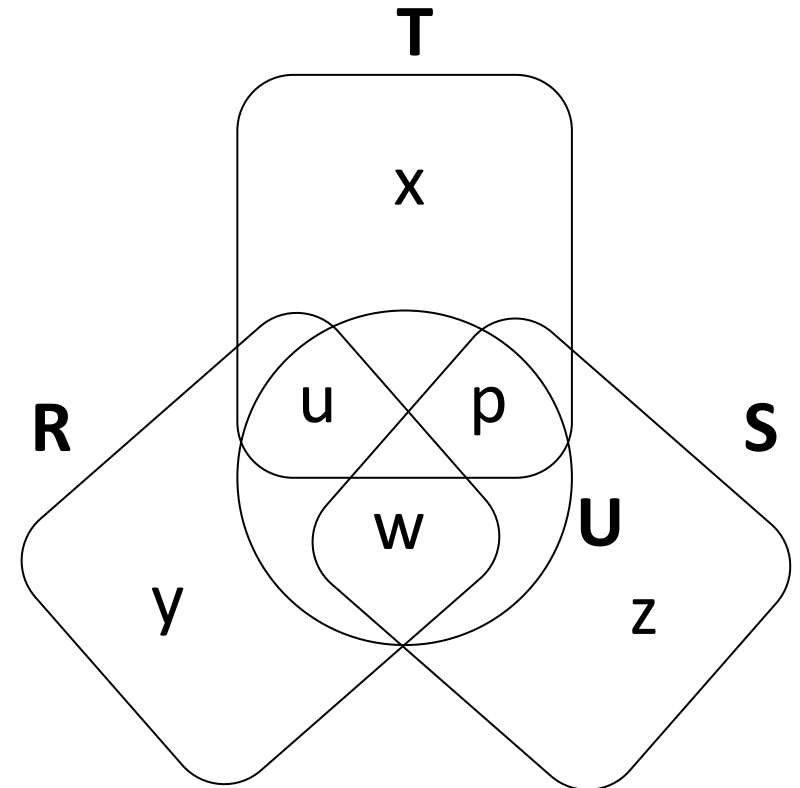
Join tree ?

$U(u,p,w)$

$R(y,u,w)$

$S(z,p,w)$

$T(x,u,p)$



GYO reduction: Example 1

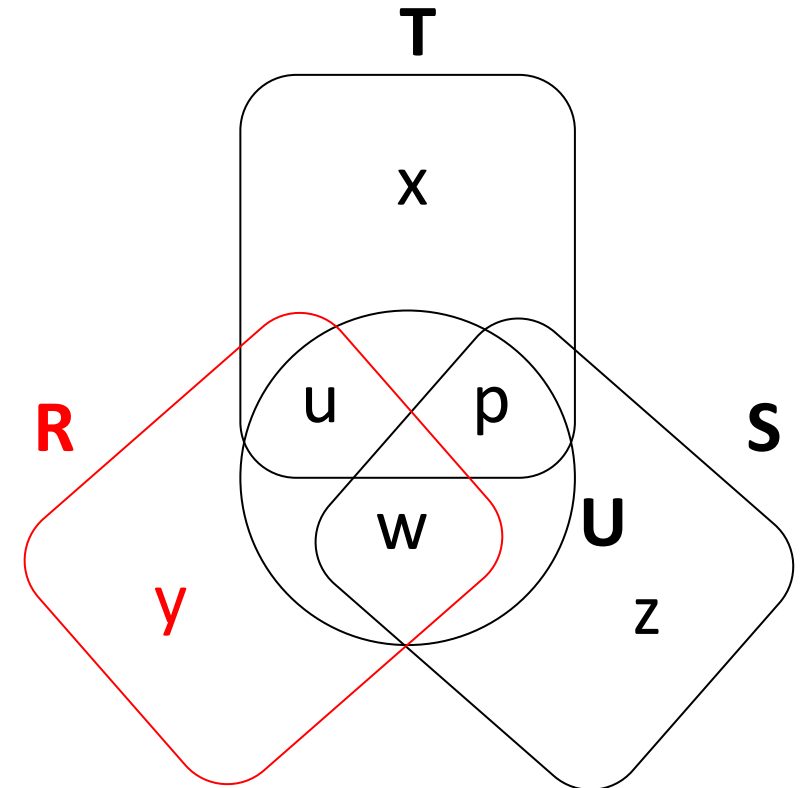
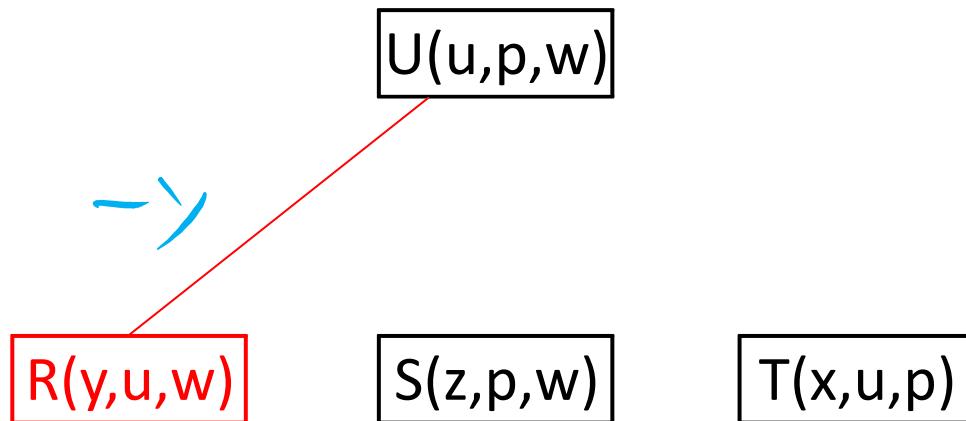


$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree



GYO reduction: Example 1

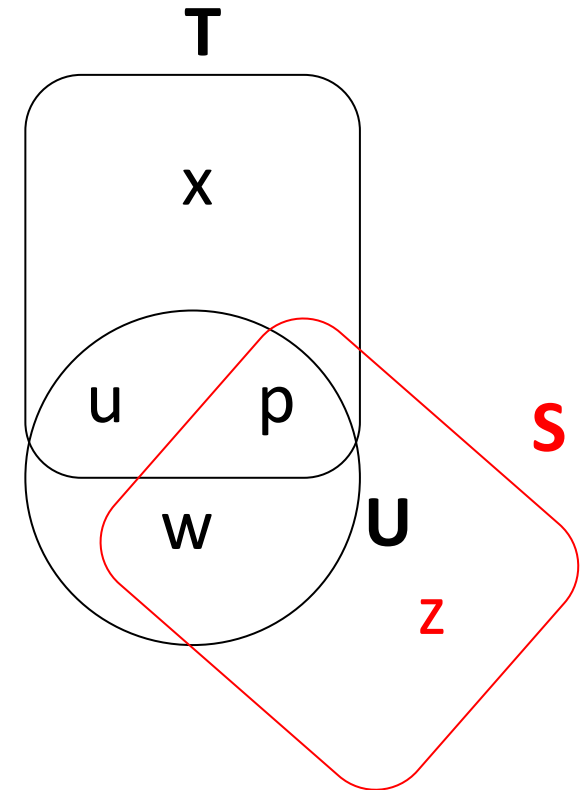
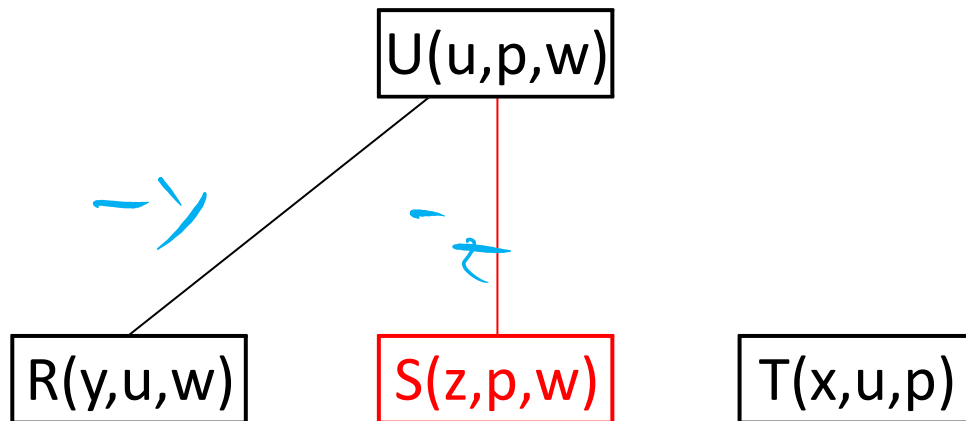


$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree



GYO reduction: Example 1

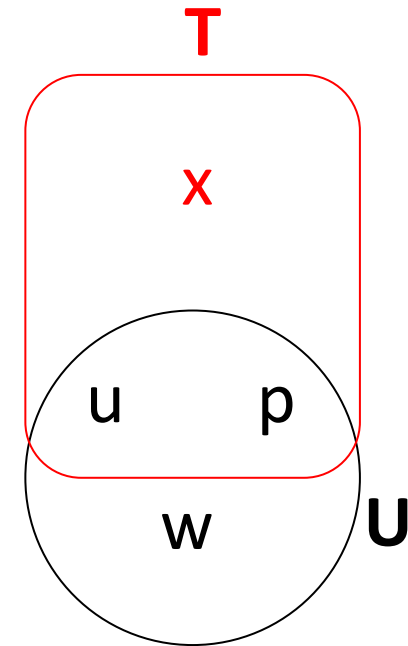
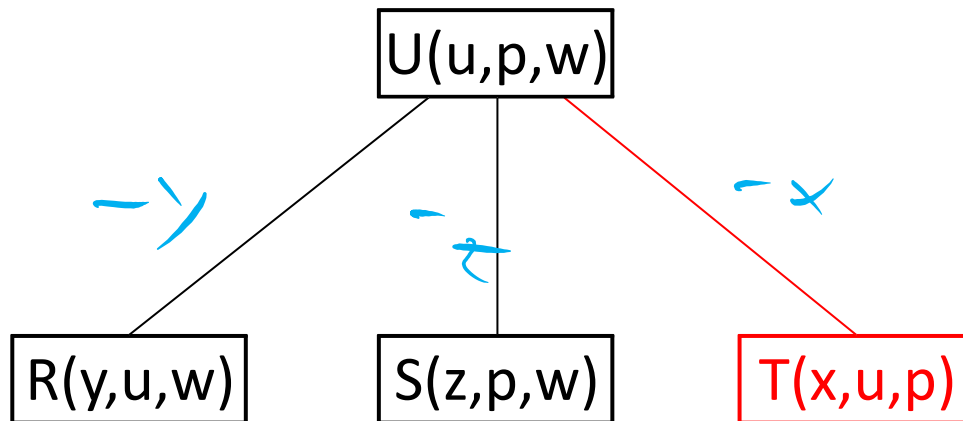


$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree



GYO reduction: Example 1

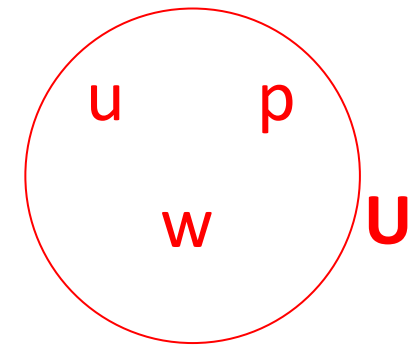
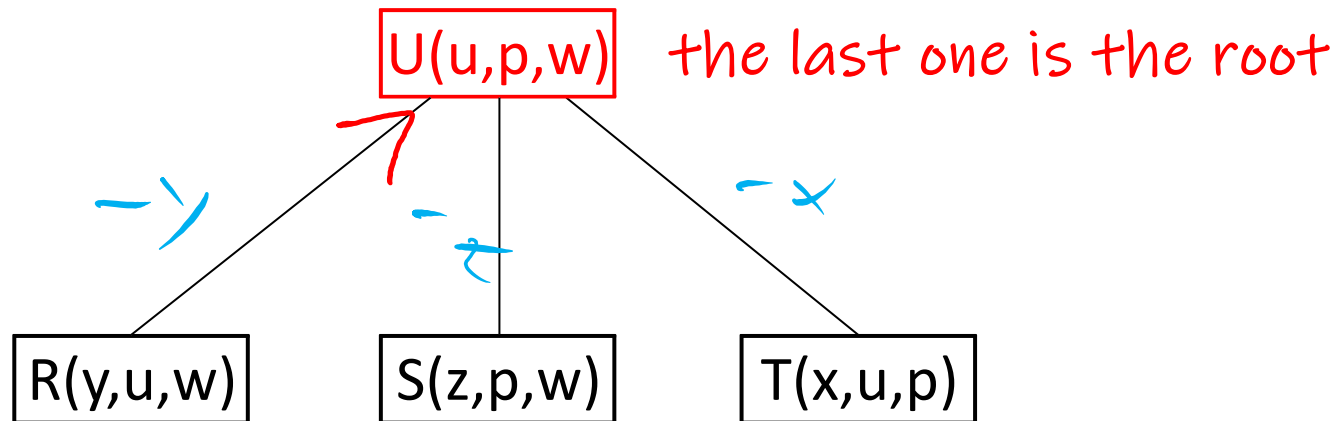


$Q :- R(y,u,w), S(z,p,w), T(x,u,p), U(u,p,w).$

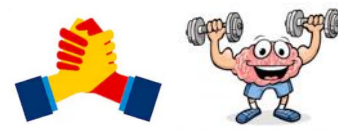
GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree



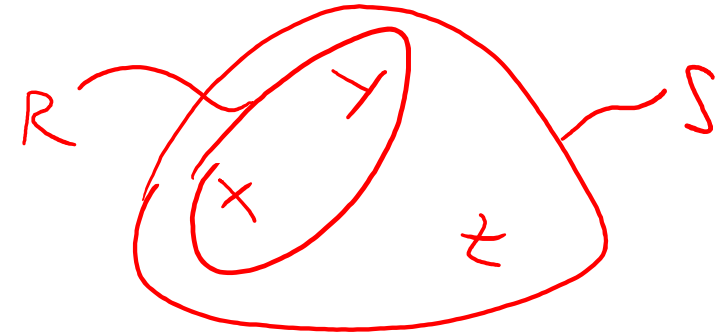
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges



Join tree

?

Query hypergraph

?

GYO reduction: Example 2



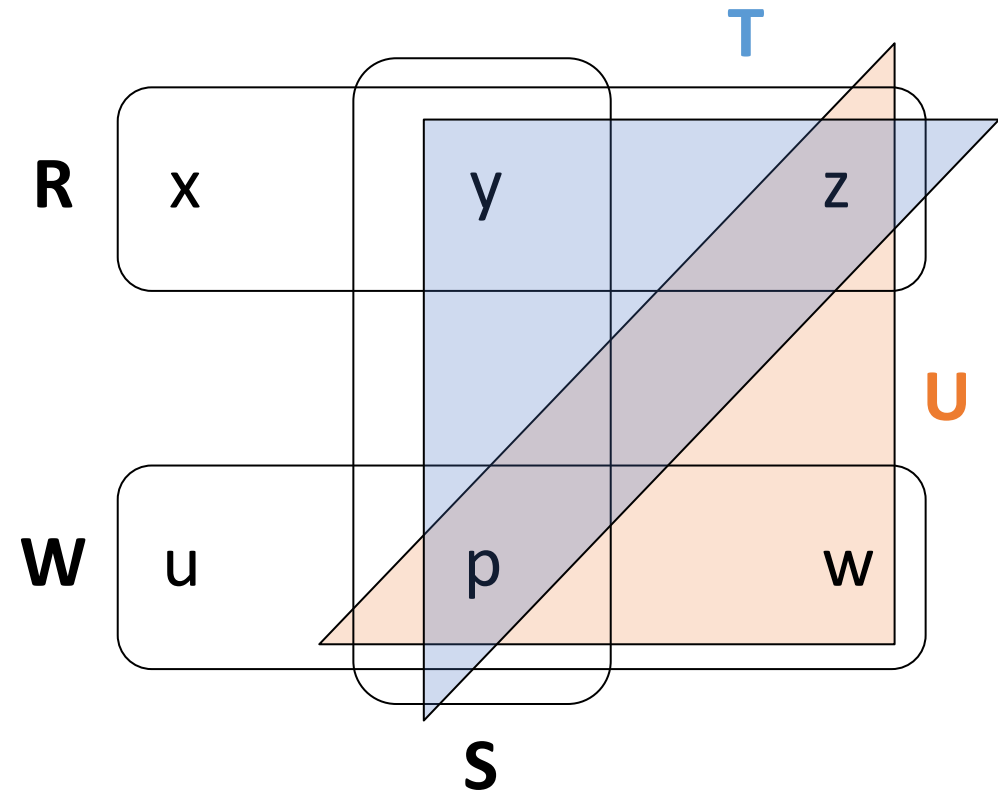
$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (**ear removal**)

- remove **isolated** nodes
- remove **consumed** or empty edges

Join tree

?



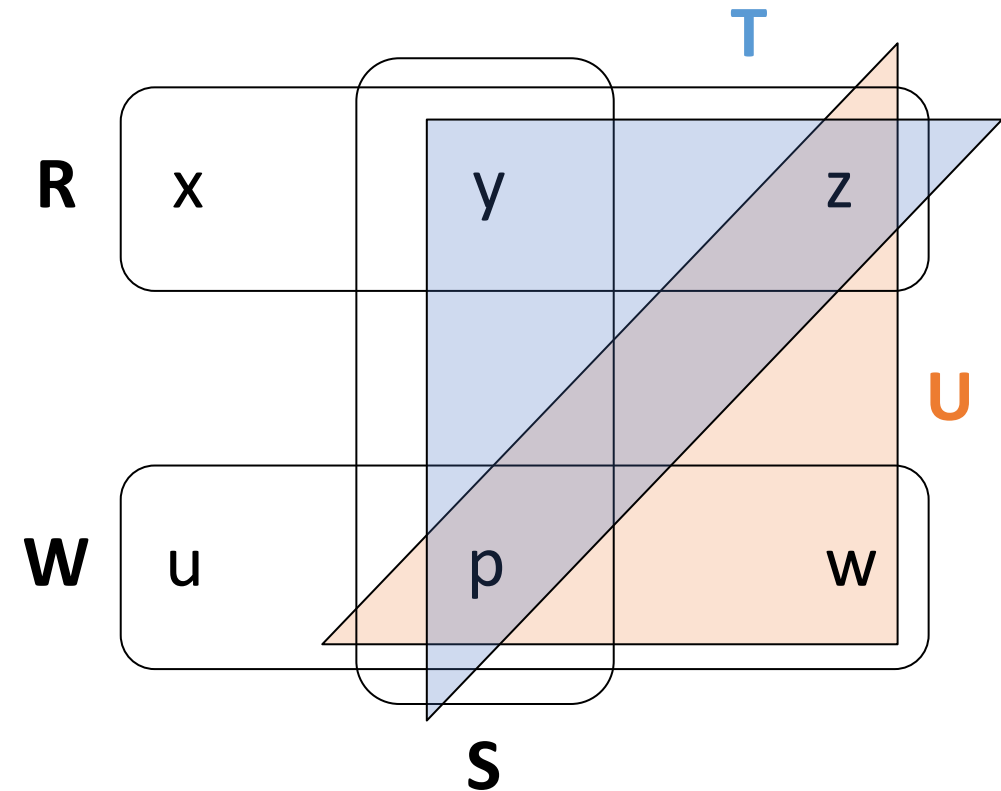
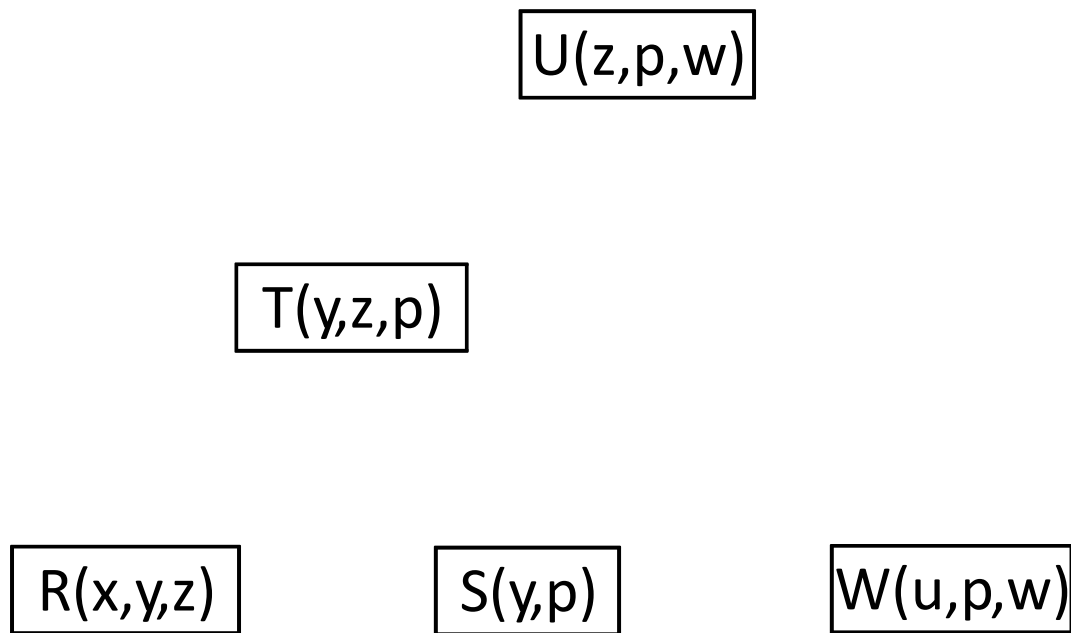
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



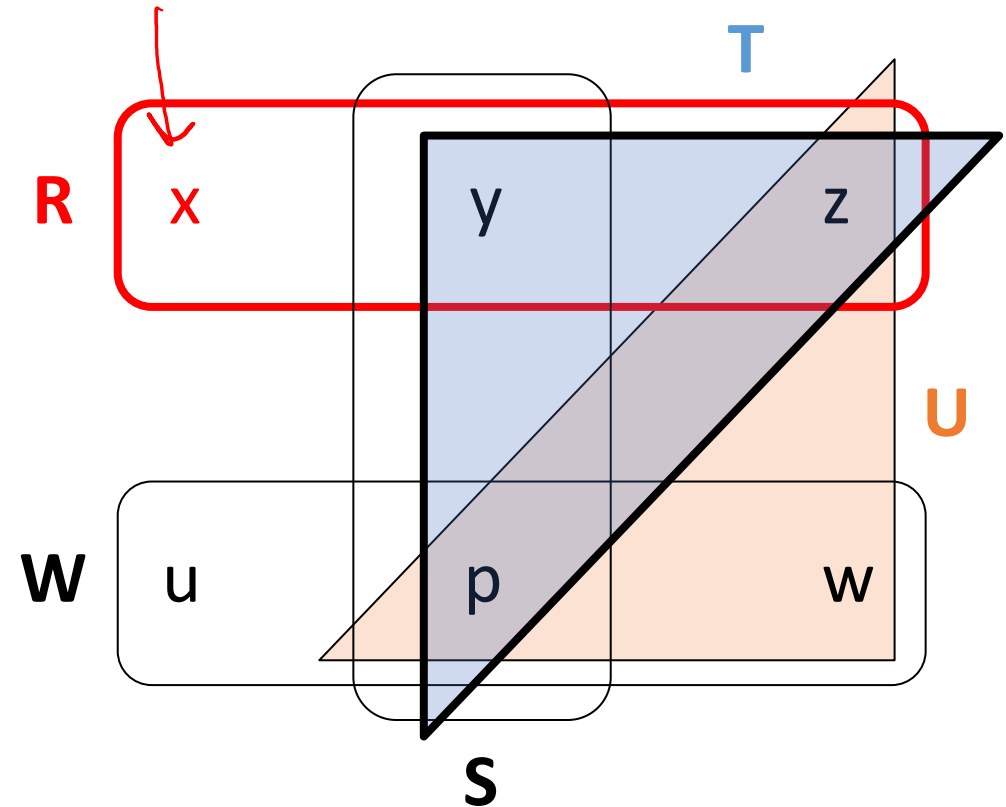
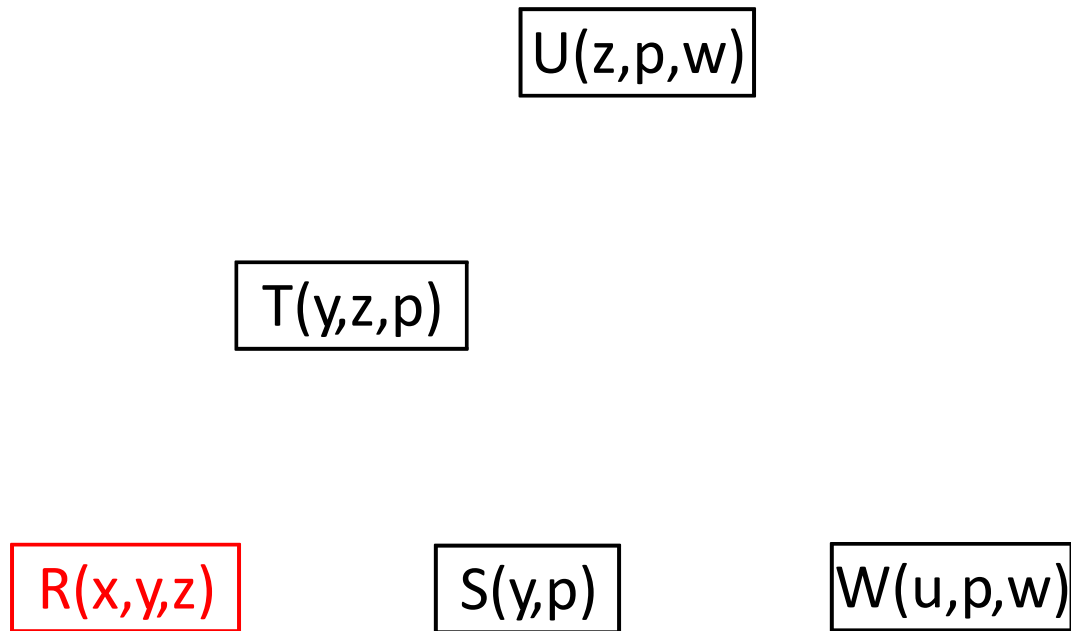
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

GYO REDUCTION (ear removal)

- remove isolated nodes
- remove consumed or empty edges



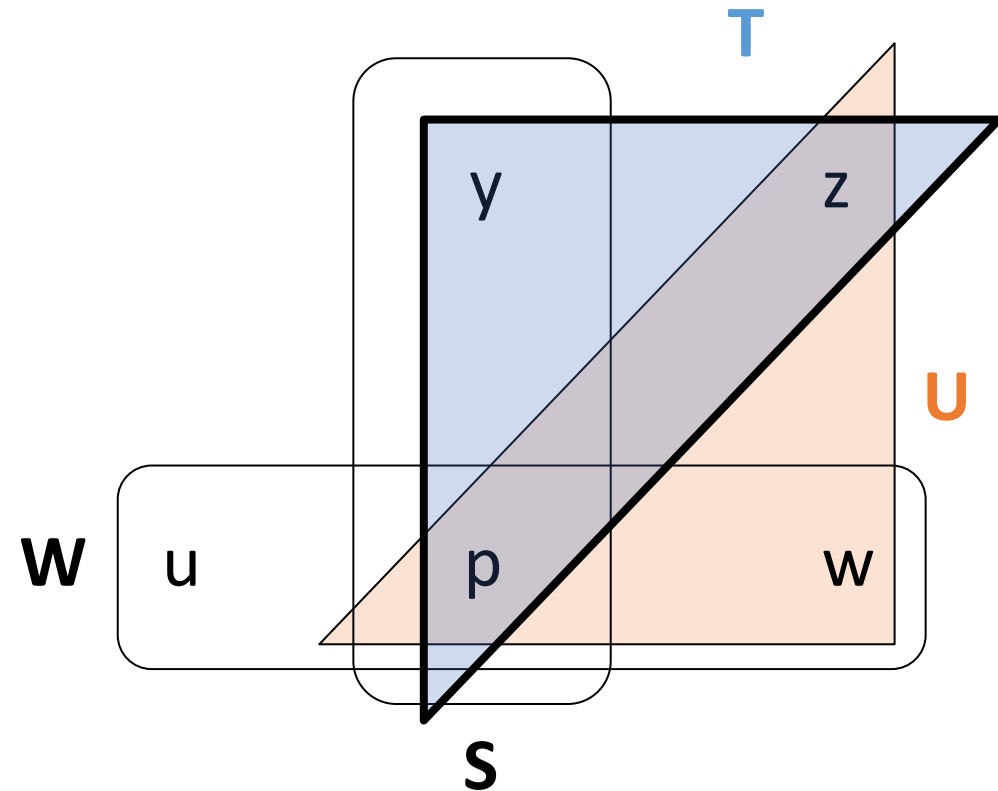
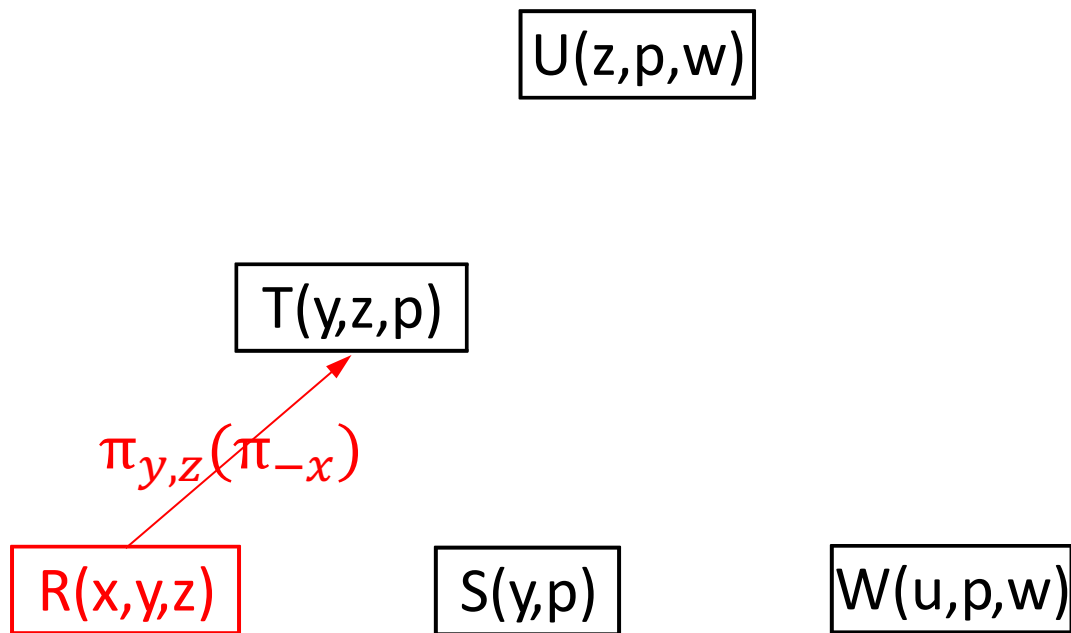
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



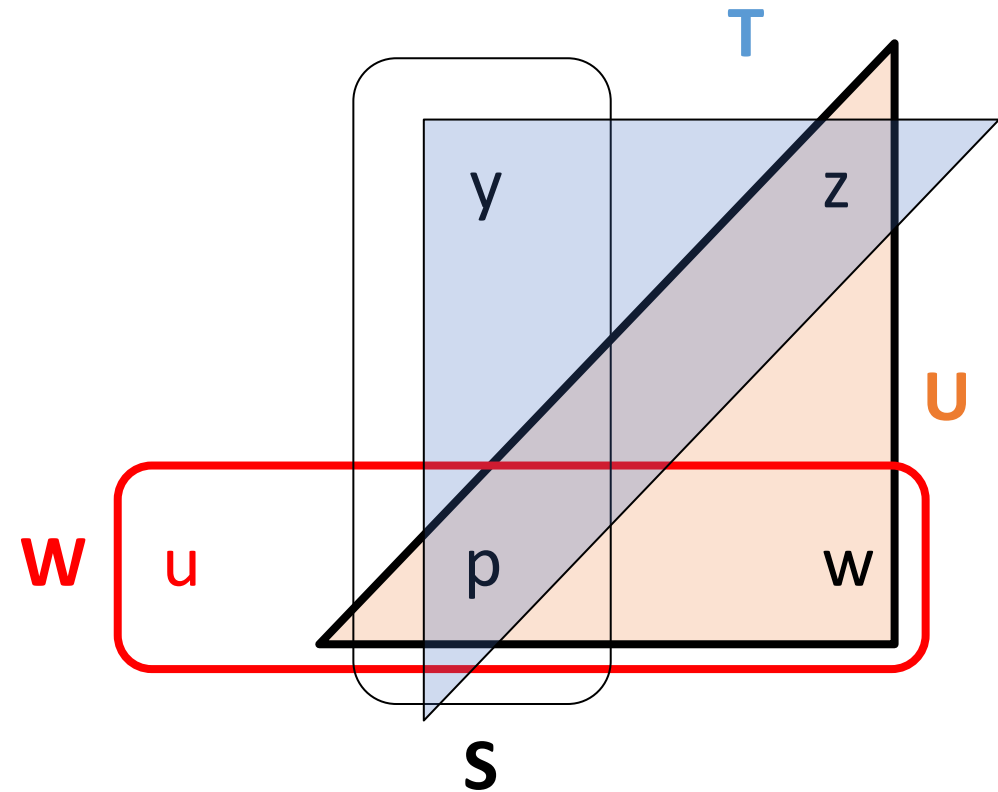
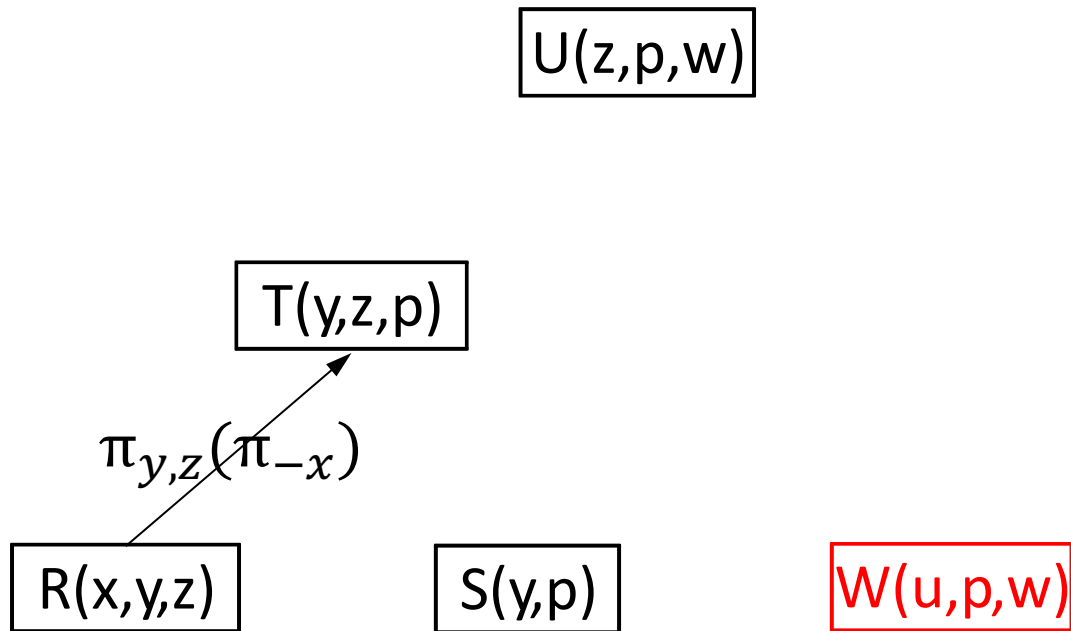
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



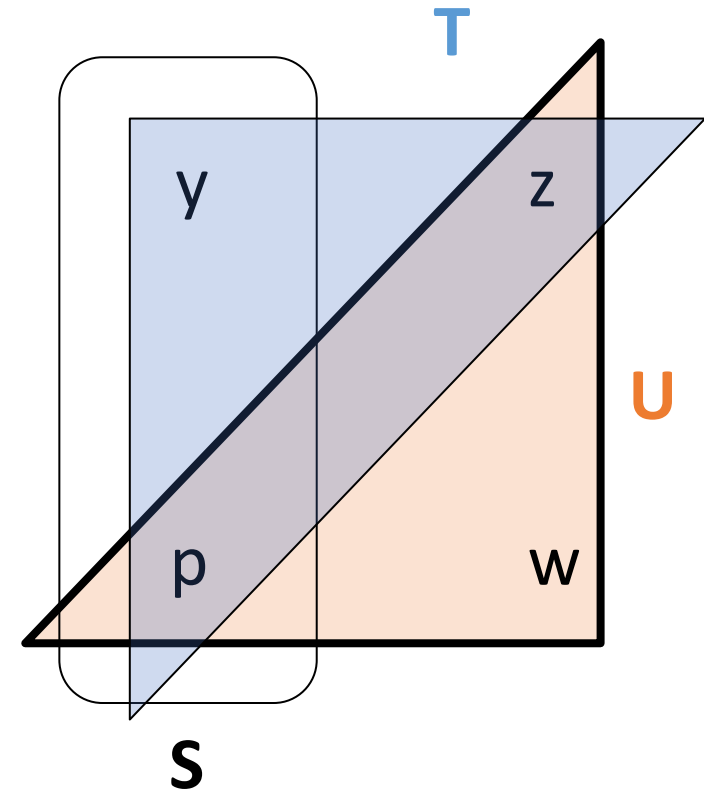
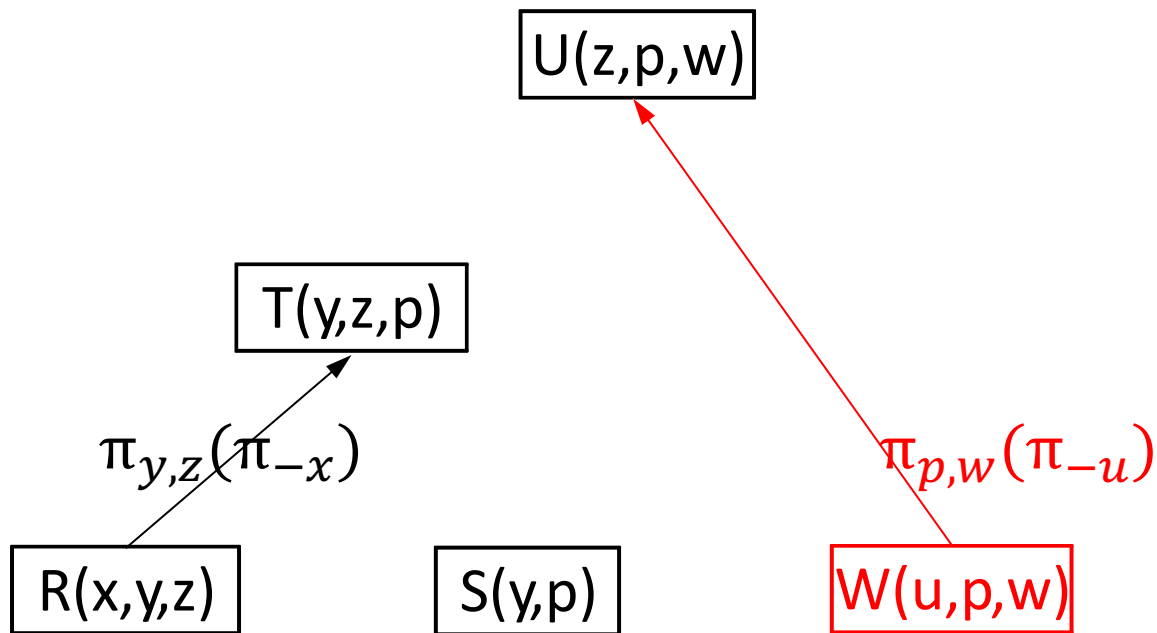
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



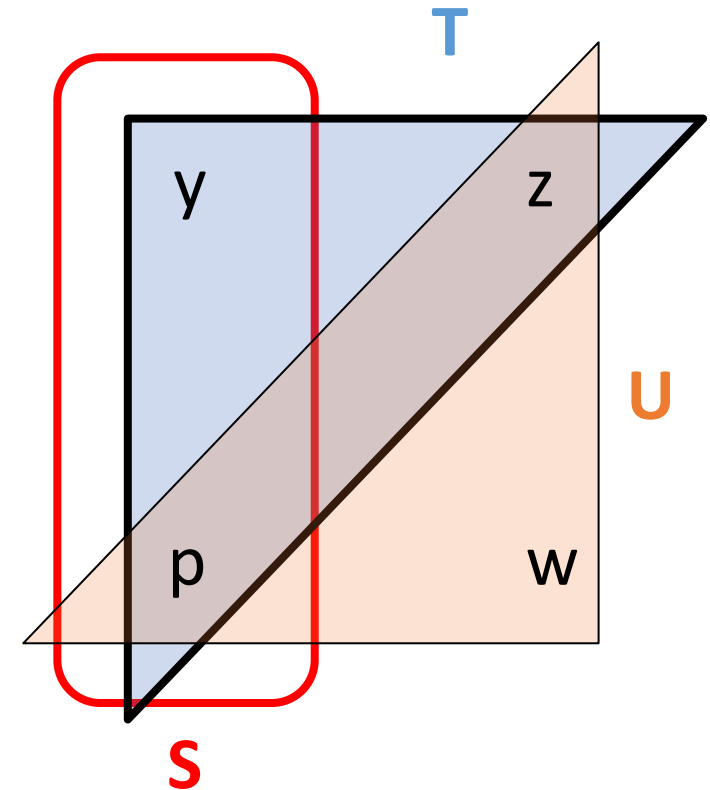
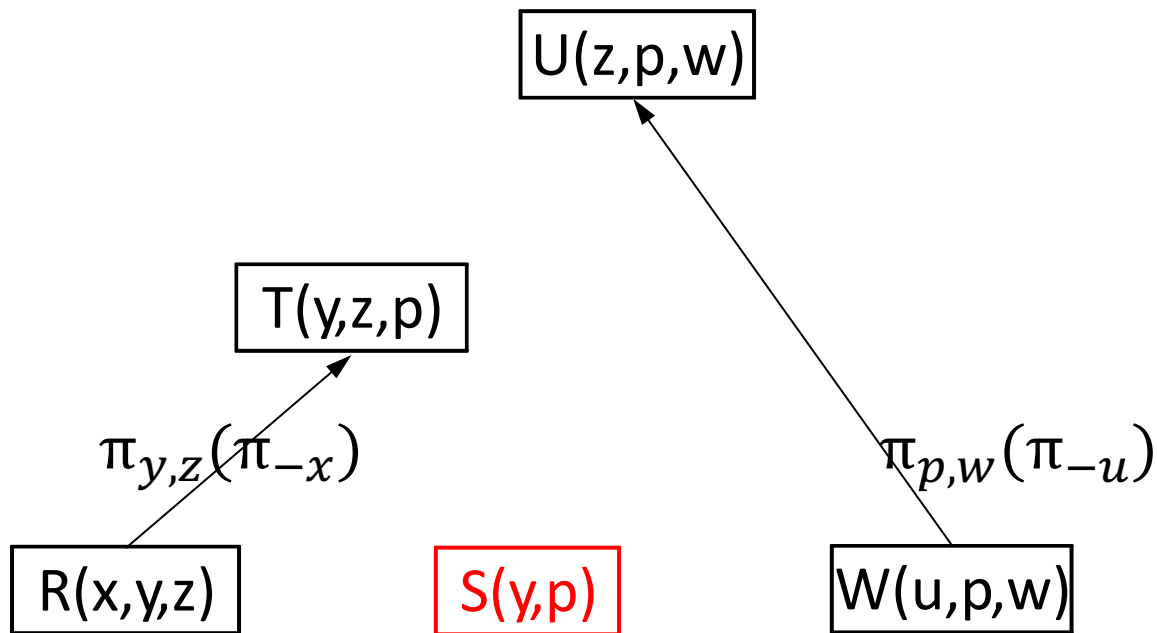
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



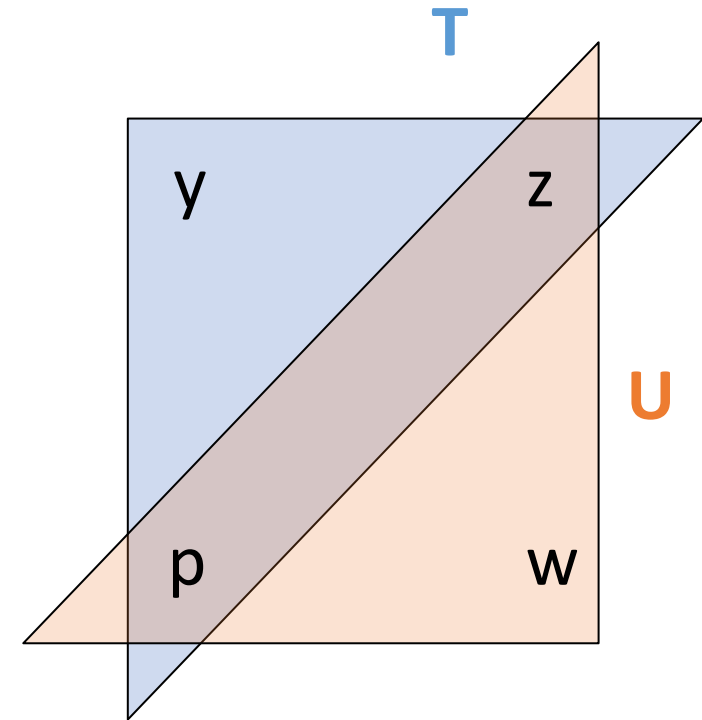
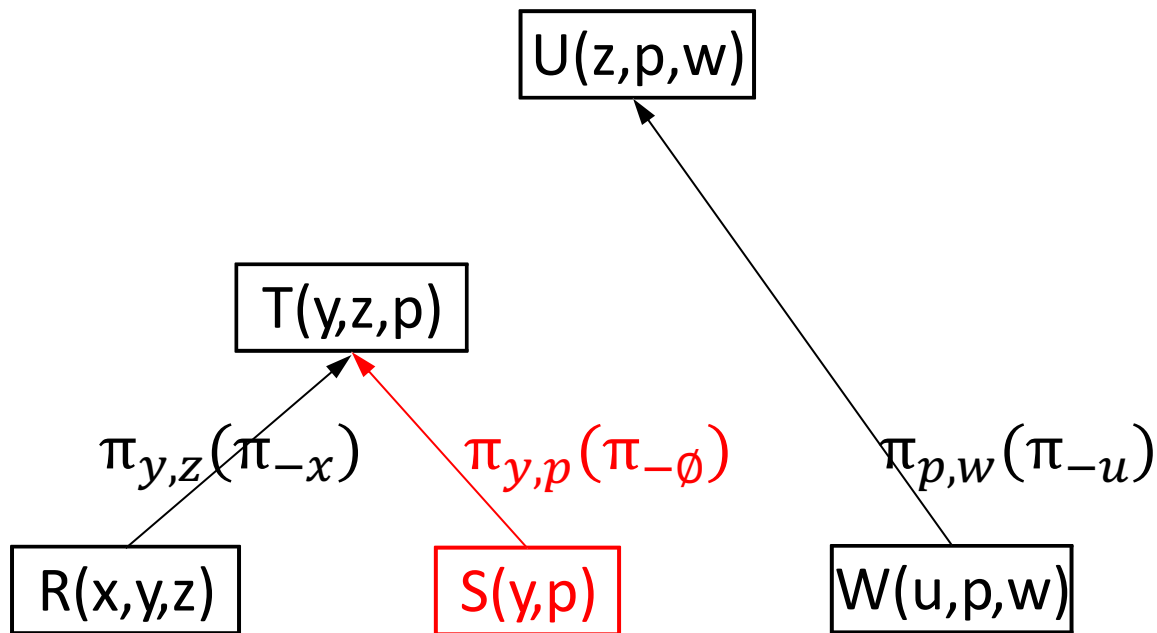
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



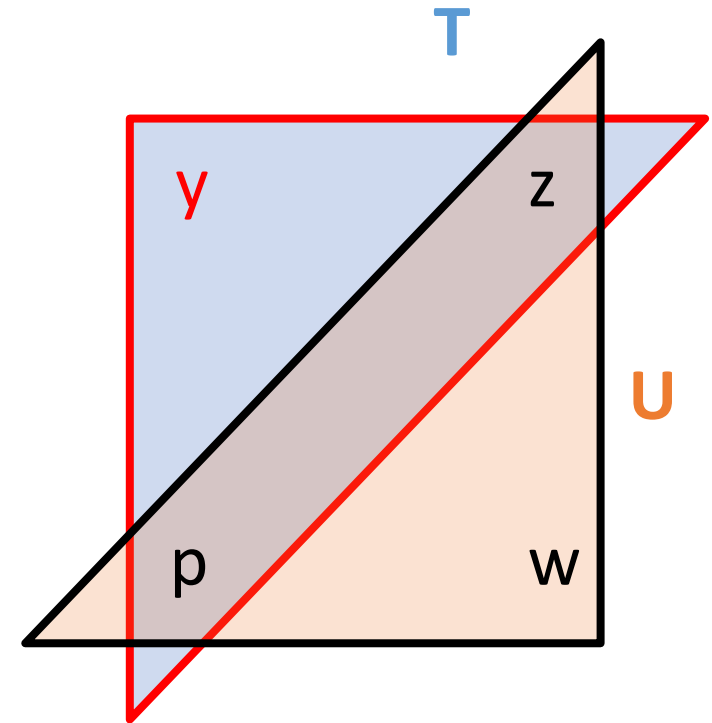
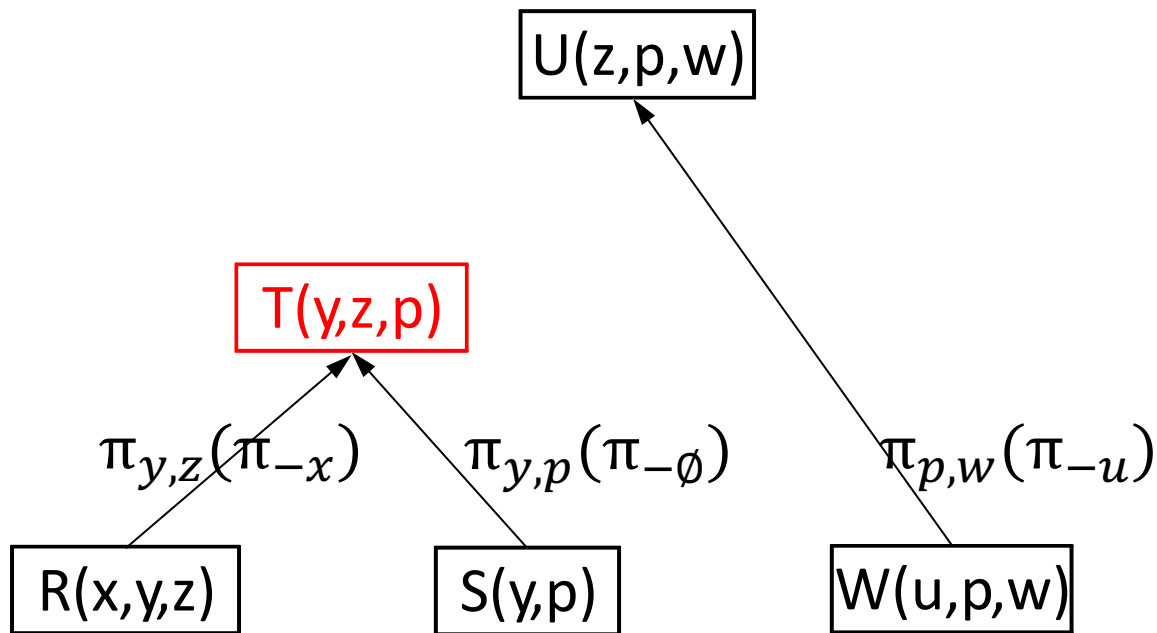
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



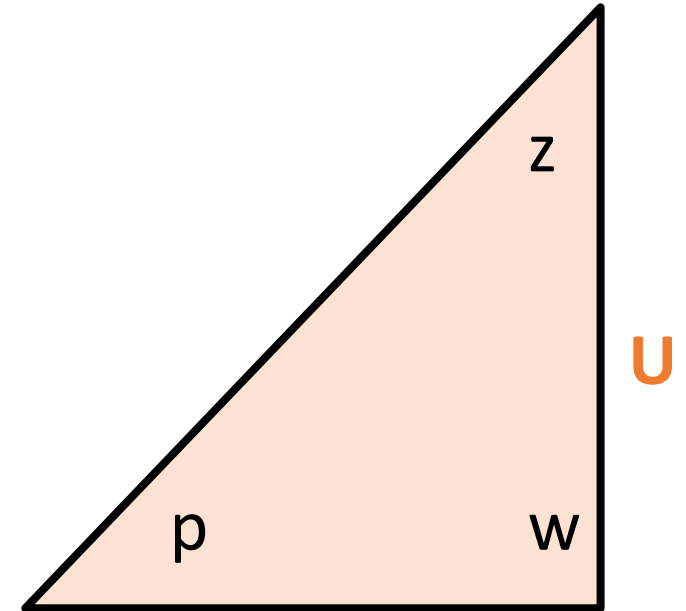
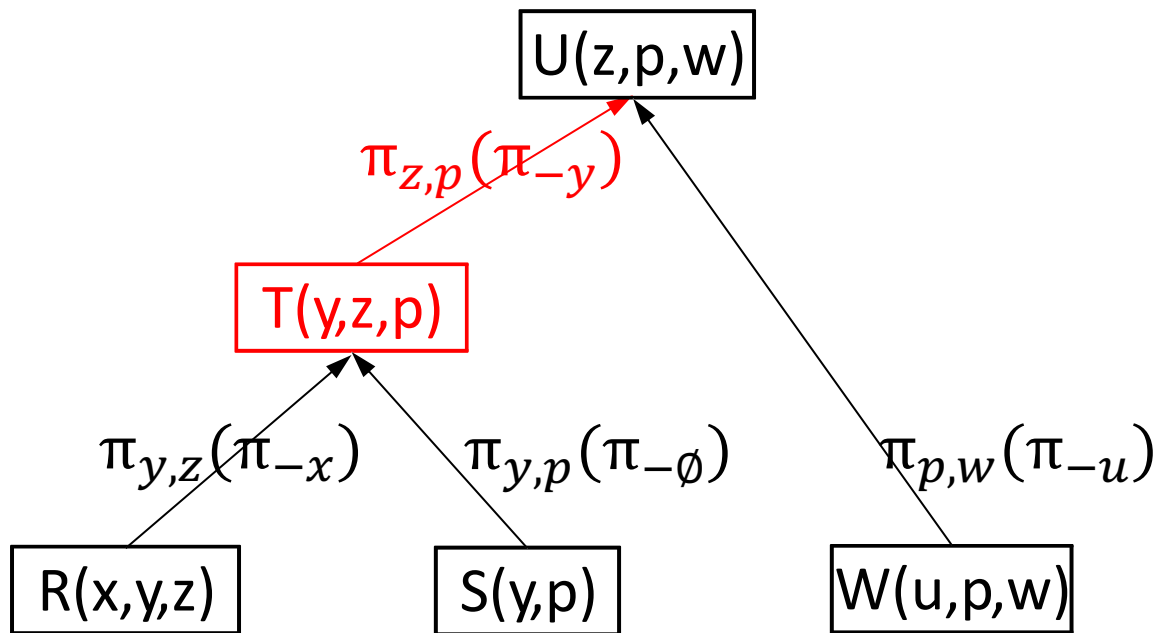
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



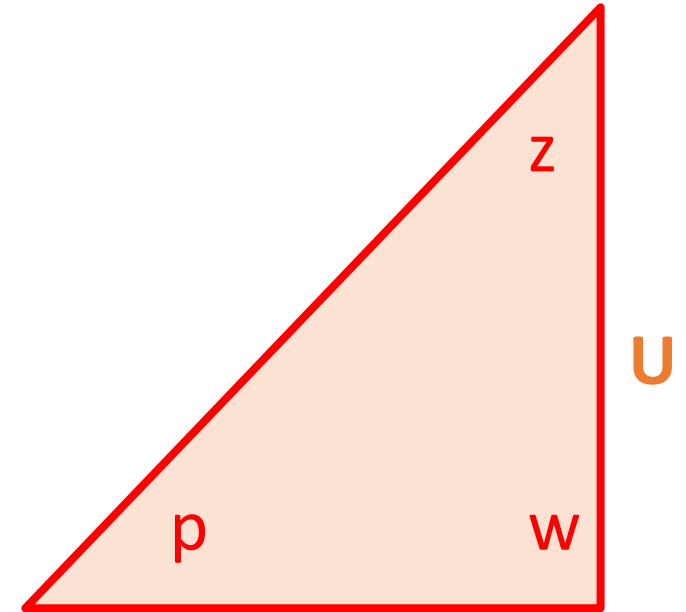
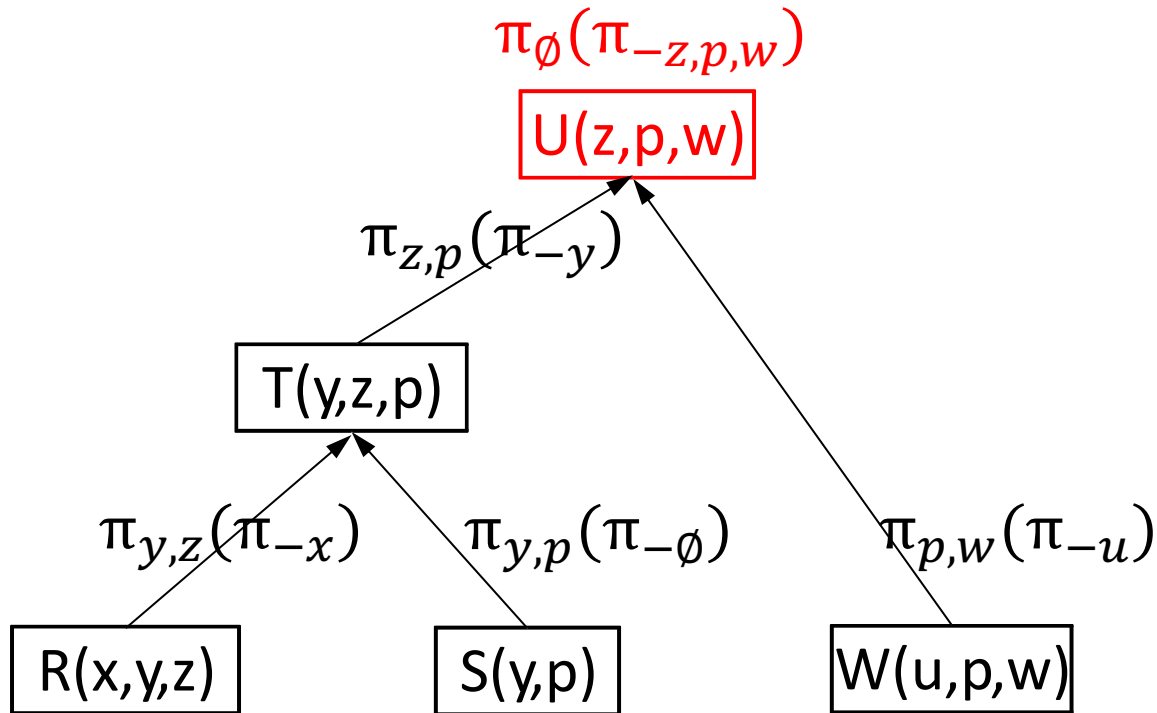
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



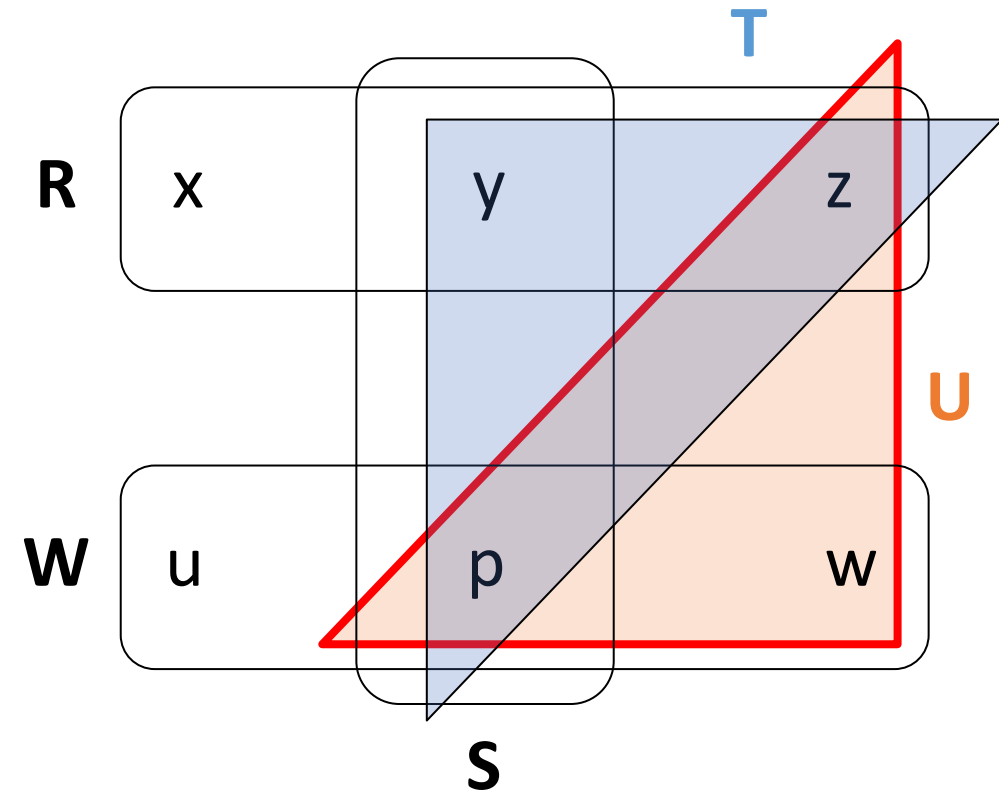
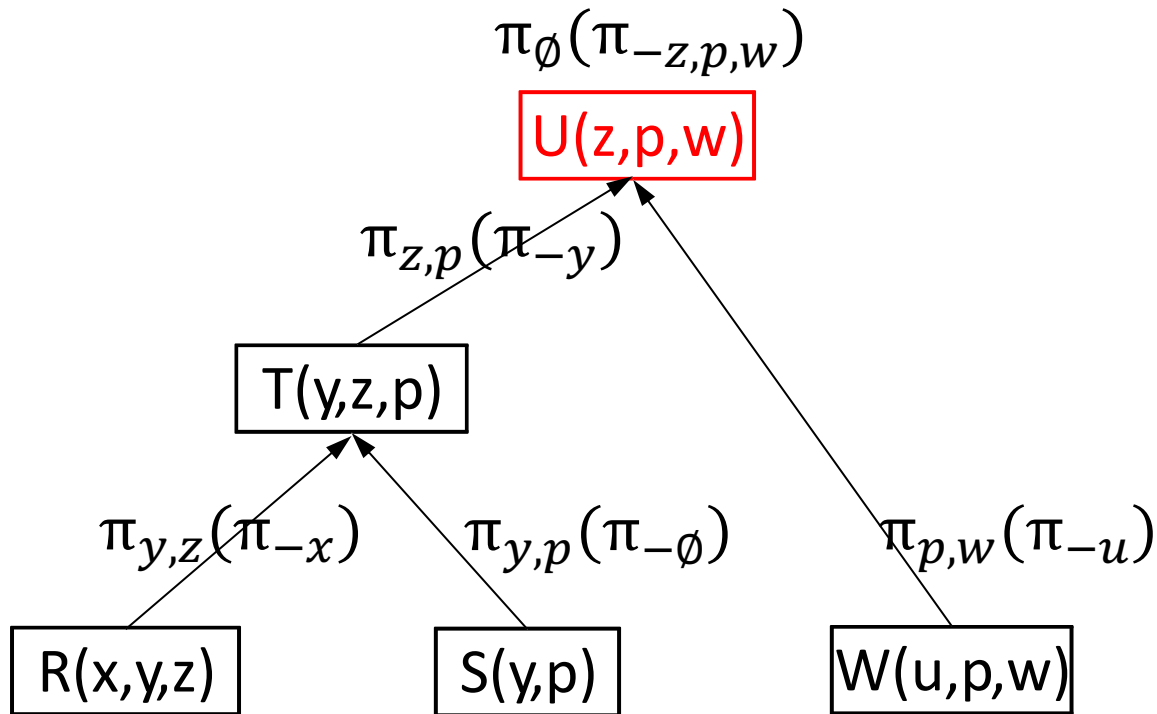
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



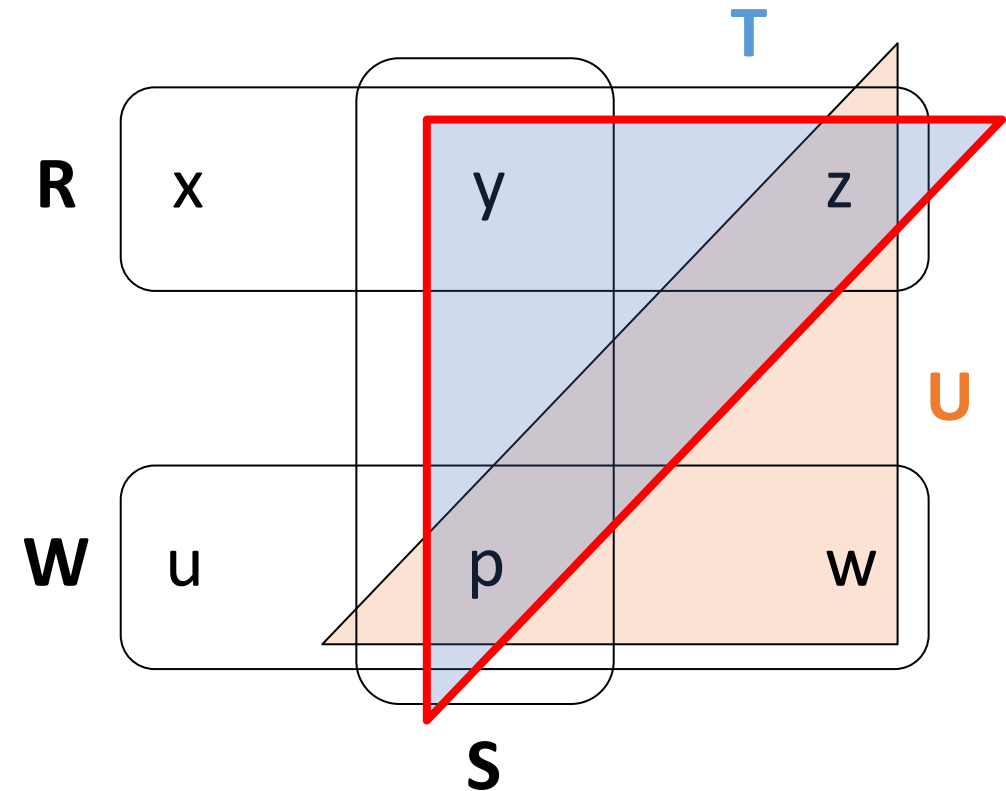
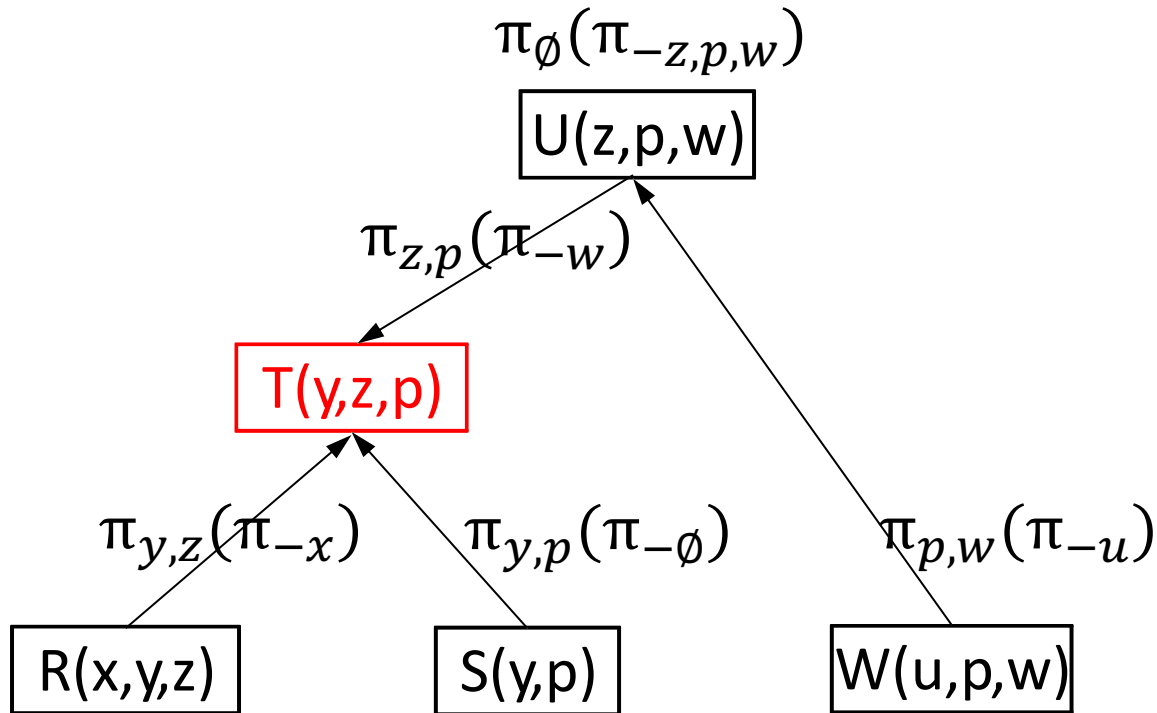
GYO reduction: Example 2



$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).$

GYO REDUCTION (ear removal)

- remove **isolated** nodes
- remove **consumed** or empty edges



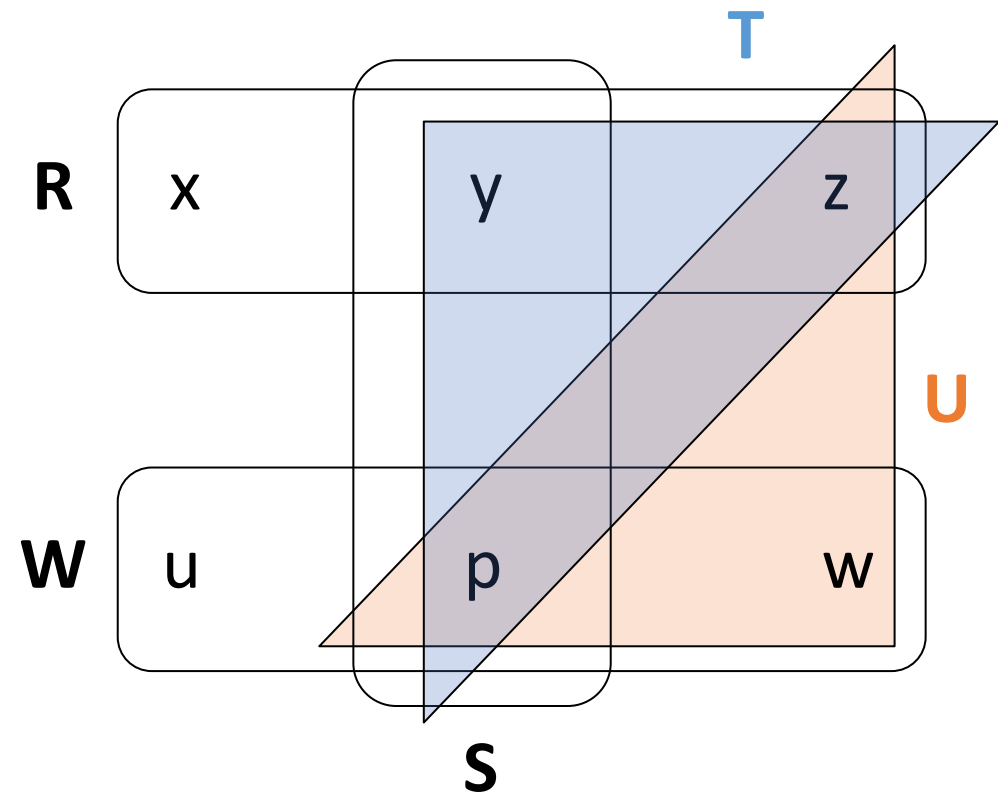
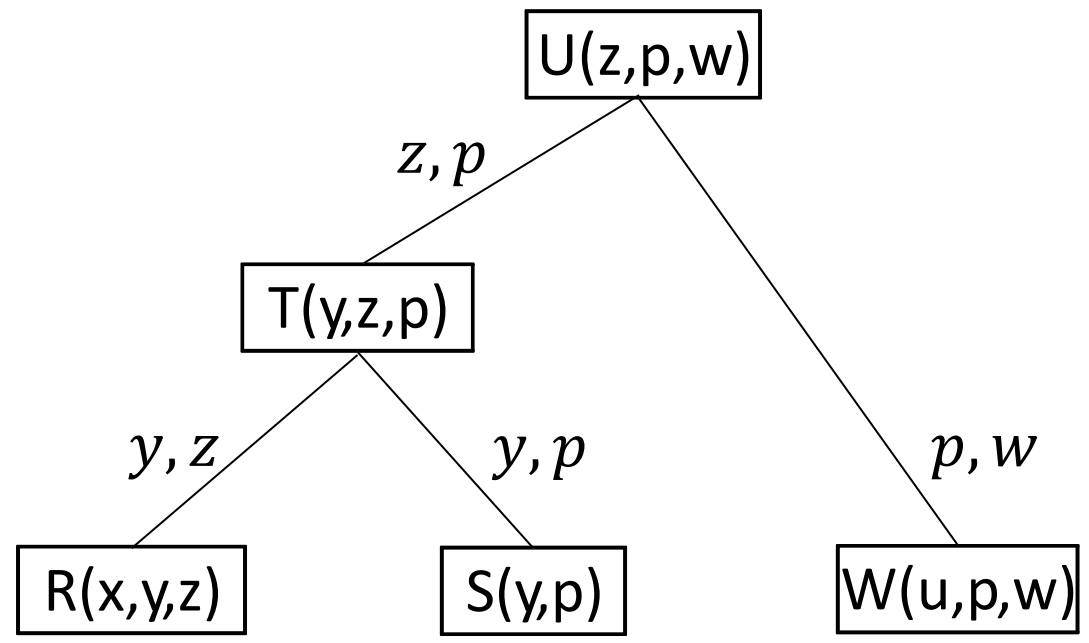
GYO reduction: Example 2



Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,u,p), W(u,p,w).

- GYO REDUCTION (ear removal)
- remove isolated nodes
 - remove consumed or empty edges

This is our join tree 😊



GYO reduction: Example 3: Are join trees unique?



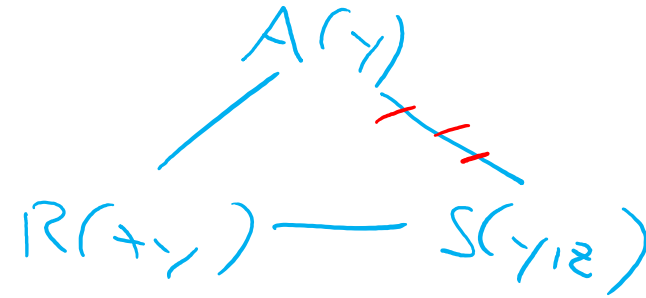
Q :- R(x,y), S(y,z), A(y).

?

GYO reduction: Example 3: Are join trees unique?

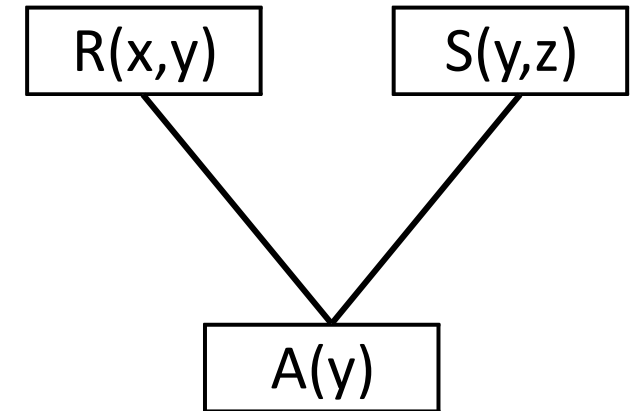
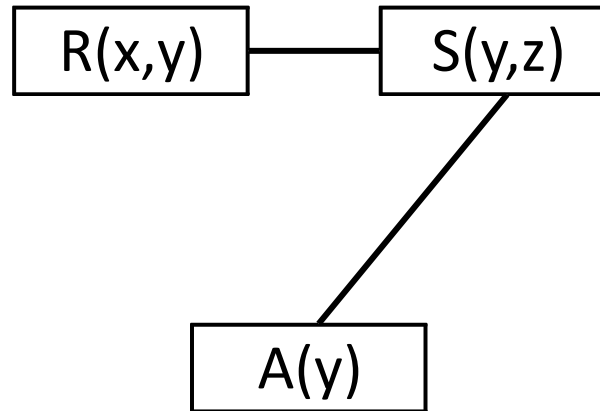
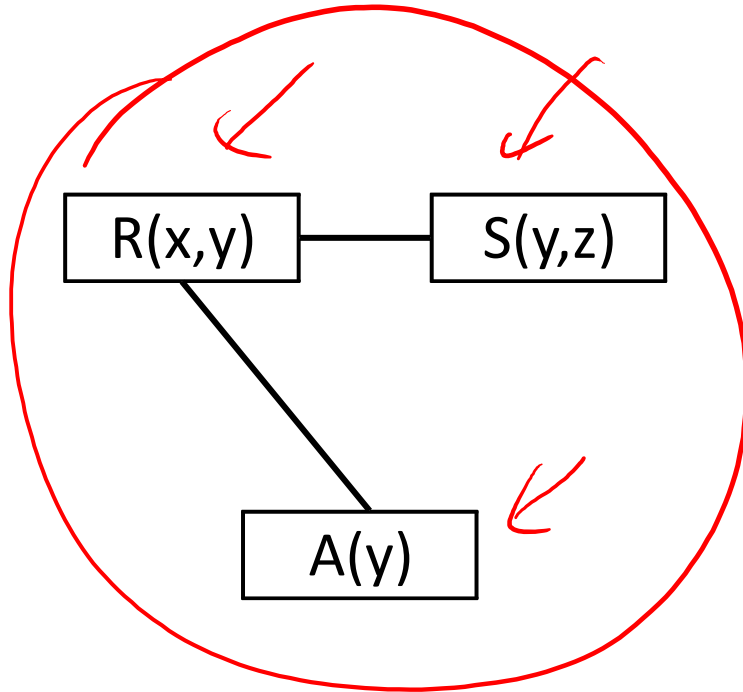


Q :- R(x,y), S(y,z), A(y).



3 CHOICES FOR ROOT

In addition, we have 3 choices as roots for each of the 3 different join trees. Leads to 9 different rooted join trees!



Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

<https://northeastern-datalab.github.io/>

Semijoin Reducer



$$Q_3^\infty(x,y,z) = R(x,y) \bowtie S(y,z) \bowtie T(z,w)$$

DEFINITION: A **full semijoin reducer** (semijoin program) is:

a sequence of semijoins on the join tree

$$R'(x,y) = R(y,z) \bowtie \dots$$

$$S'(y,z) = S(y,z) \bowtie \dots$$

...

s.t. there no more "dangling tuples" in the reduced relations

Then you can rewrite the query over the reduced relations

$$Q(x,y,z) = R'(x,y) \bowtie S'(y,z) \bowtie T'(z,w)$$

Semijoin Reducer



$$Q_3^\infty(x, y, z) = R(x, y) \bowtie S(y, z) \bowtie T(z, w)$$

A full reducer is

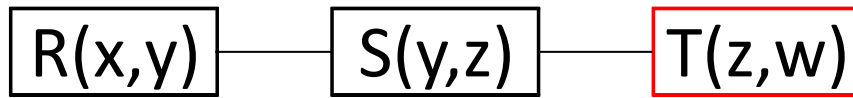
?

Semijoin Reducer



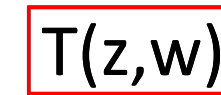
$$Q_3^\infty(x,y,z) = R(x,y) \bowtie S(y,z) \bowtie T(z,w)$$

A full reducer is

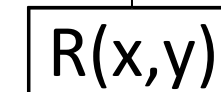
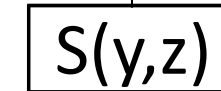


?

1. Find a join tree



2. pick a root

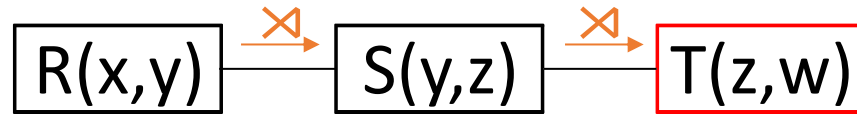


Semijoin Reducer



$$Q_3^\infty(x,y,z) = R(x,y) \bowtie S(y,z) \bowtie T(z,w)$$

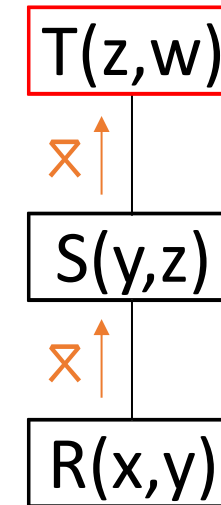
A full reducer is



3. collect at root (= bottom-up)

$$S_1(y,z) = S(y,z) \bowtie R(x,y)$$
$$T_1(z,w) = T(z,w) \bowtie S_1(y,z)$$

1. Find a join tree



2. pick a root

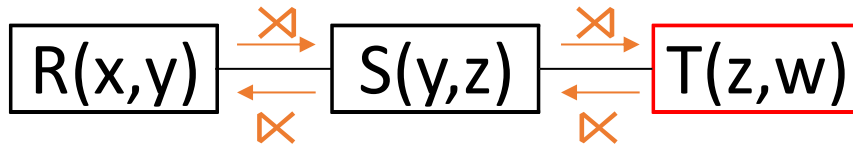
3. up

Semijoin Reducer



$$Q_3^\infty(x,y,z) = R(x,y) \bowtie S(y,z) \bowtie T(z,w)$$

A full reducer is



3. collect at root (= bottom-up)

$$S_1(y,z) = S(y,z) \bowtie R(x,y)$$

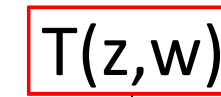
$$T_1(z,w) = T(z,w) \bowtie S_1(y,z)$$

4. distribute to leaves (= top-down)

$$S_2(z,y) = S_1(y,z) \bowtie T_1(z,y)$$

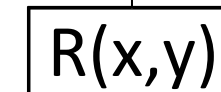
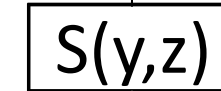
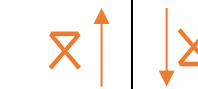
$$R_1(x,y) = R(x,y) \bowtie S_2(y,z)$$

1. Find a join tree



2. pick a root

3. up



4. down

5. The rewritten query is

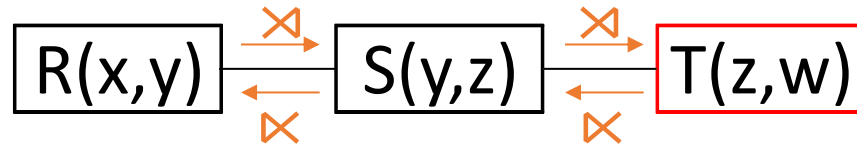
?

Semijoin Reducer



$$Q_3^\infty(x,y,z) = R(x,y) \bowtie S(y,z) \bowtie T(z,w)$$

A full reducer is



3. collect at root (= bottom-up)

$$S_1(y,z) = S(y,z) \bowtie R(x,y)$$

$$T_1(z,w) = T(z,w) \bowtie S_1(y,z)$$

4. distribute to leaves

$$S_2(z,y) = S_1(y,z) \bowtie T_1(z,y)$$

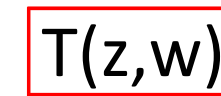
(= top-down)

$$R_1(x,y) = R(x,y) \bowtie S_2(y,z)$$

5. The rewritten query is

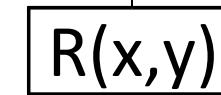
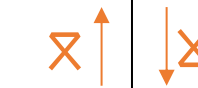
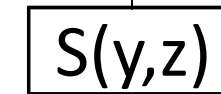
$$Q(x,y,z) = R_1(x,y) \bowtie S_2(y,z) \bowtie T_1(z,w)$$

1. Find a join tree



2. pick a root

3. up



4. down

Notice that with $2(k-1)$ messages, every of the k tables has received information from every other table!

Application

$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$

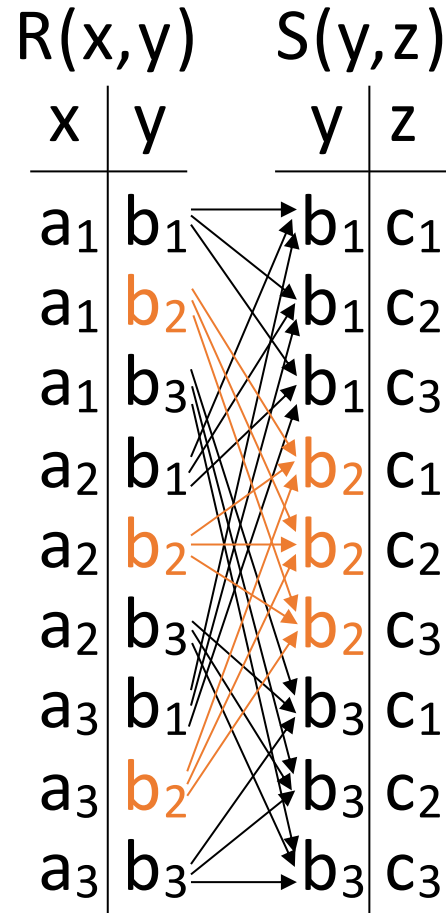
R(x,y)		S(y,z)	
x	y	y	z
a ₁	b ₁	b ₁	c ₁
a ₁	b ₂	b ₁	c ₂
a ₁	b ₃	b ₁	c ₃
a ₂	b ₁	b ₂	c ₁
a ₂	b ₂	b ₂	c ₂
a ₂	b ₃	b ₂	c ₃
a ₃	b ₁	b ₃	c ₁
a ₃	b ₂	b ₃	c ₂
a ₃	b ₃	b ₃	c ₃

How many joins are between R and S?

?

Application

$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$



How many joins are between R and S?

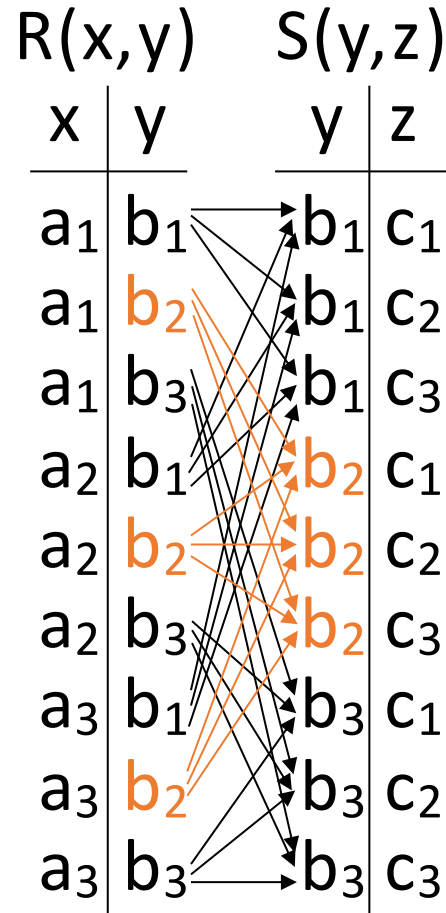
27

How can we represent the joins in a graph more compactly?

?

Application

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$$



How many joins are between R and S?

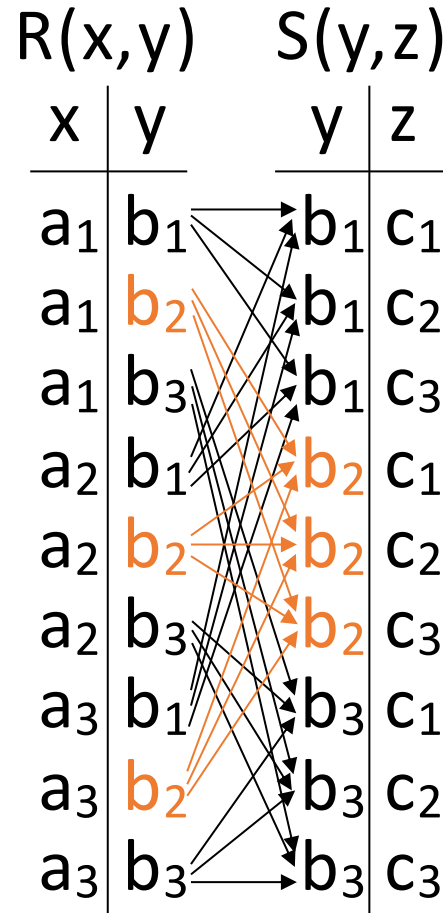
27

How can we represent the joins in a graph more compactly?

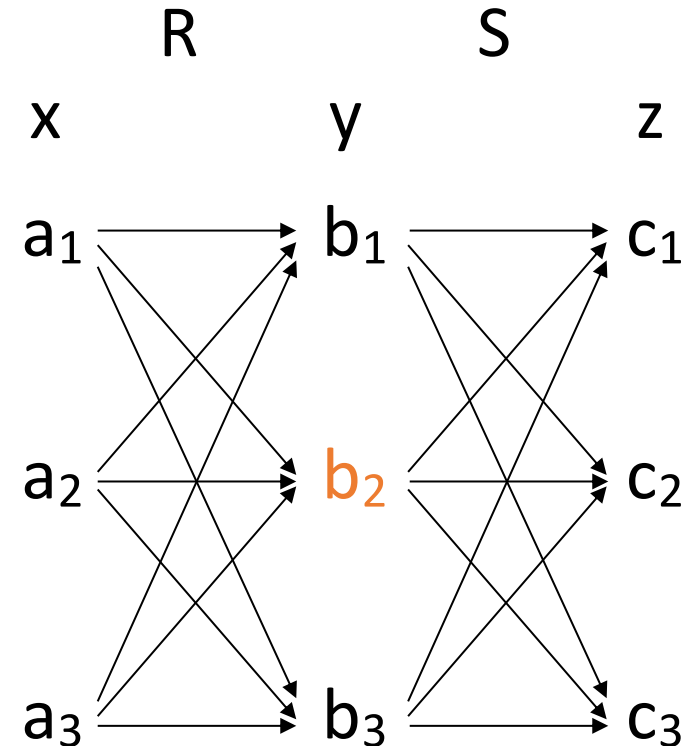
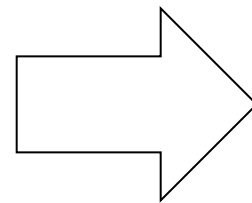
By using "the dual" (nodes as joins = domain values) and counting paths!

Application

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$$



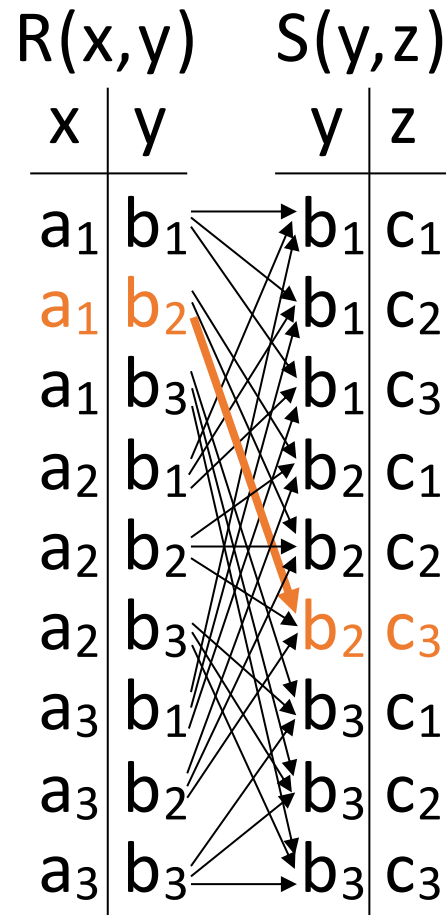
- Nodes are the tuples
- Edges are the joins



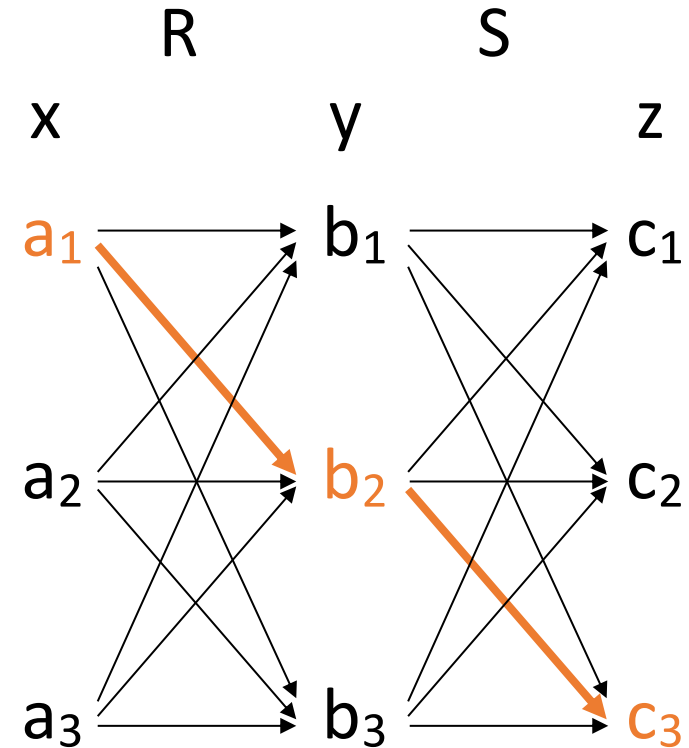
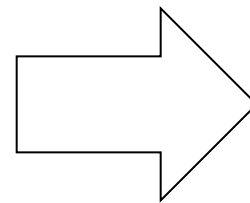
- Nodes are domain values
- Edges are the tuples
- Equi-joins b/w tuples as shared domain

Application

$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$



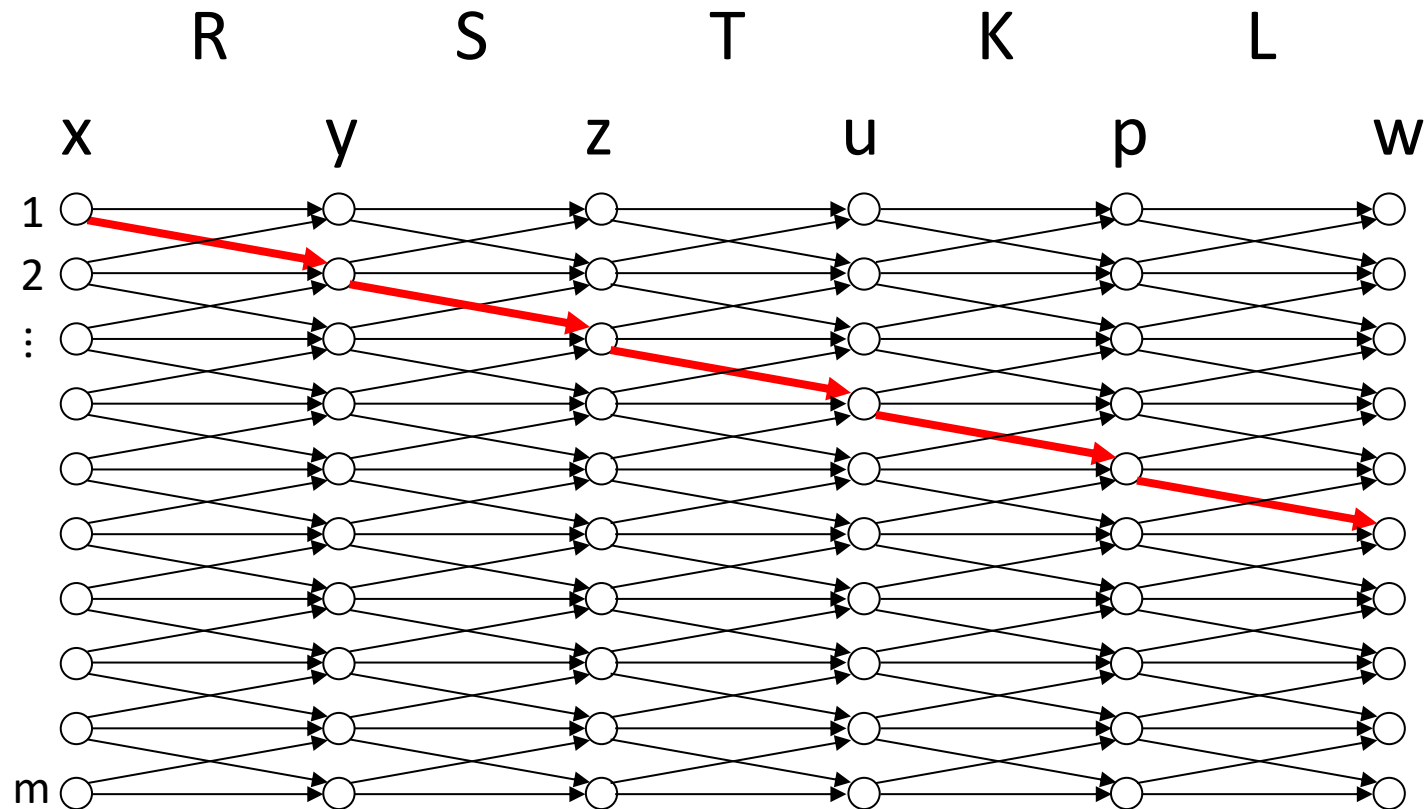
- Nodes are the tuples
- Edges are the joins



- Nodes are domain values
- Edges are the tuples
- Equi-joins b/w tuples as shared domain

Application

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y),S(y,z),T(z,u),K(u,v),L(v,w).$$

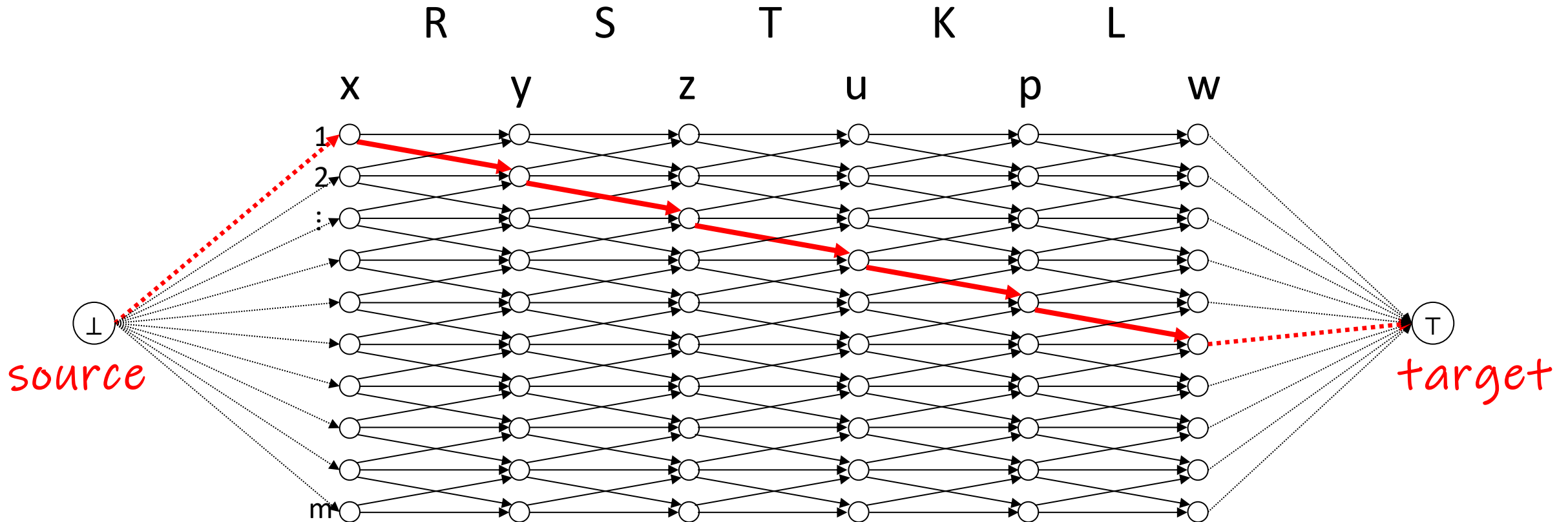


- Intermediate relations of size up to m^5 (m is here domain size!)
- But final answer can be as small as 0 (or 1...)

Application

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$$

Each source-target path corresponds to a join (this is unrelated to the flow Min-Cut-Max-Flow algorithms discussed earlier)

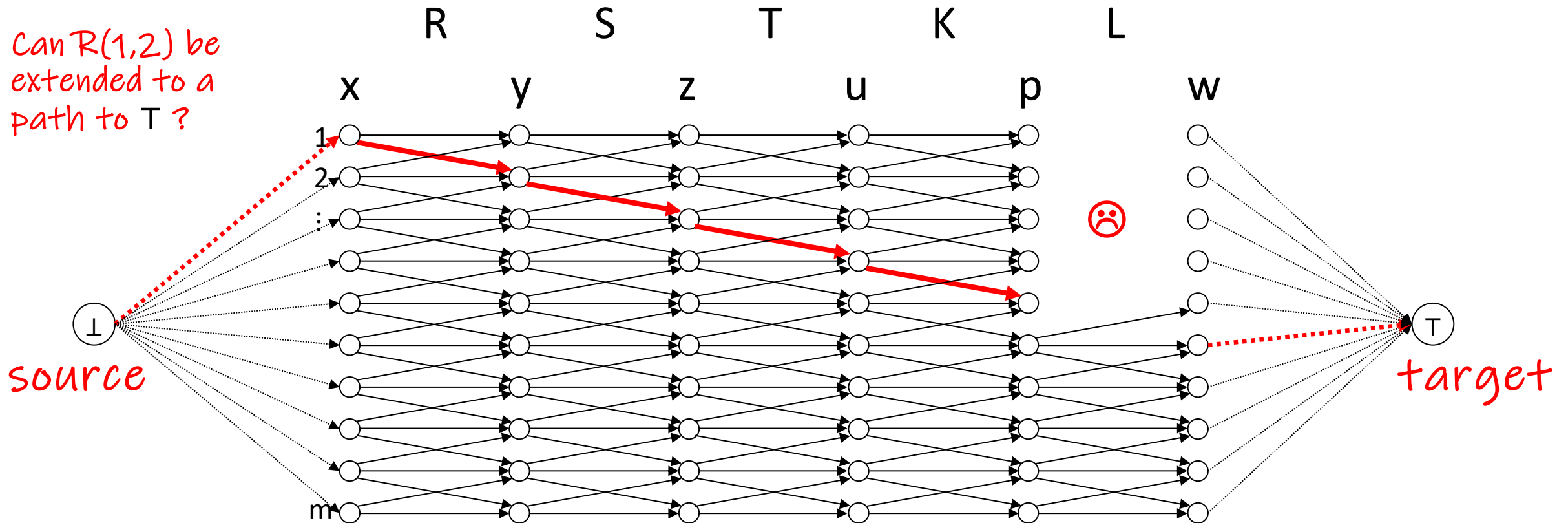


- Intermediate relations of size up to m^5 (m is here domain size!)
- But final answer can be as small as 0 (or 1...)

Application

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$$

Can $R(1,2)$ be extended to a path to T ?



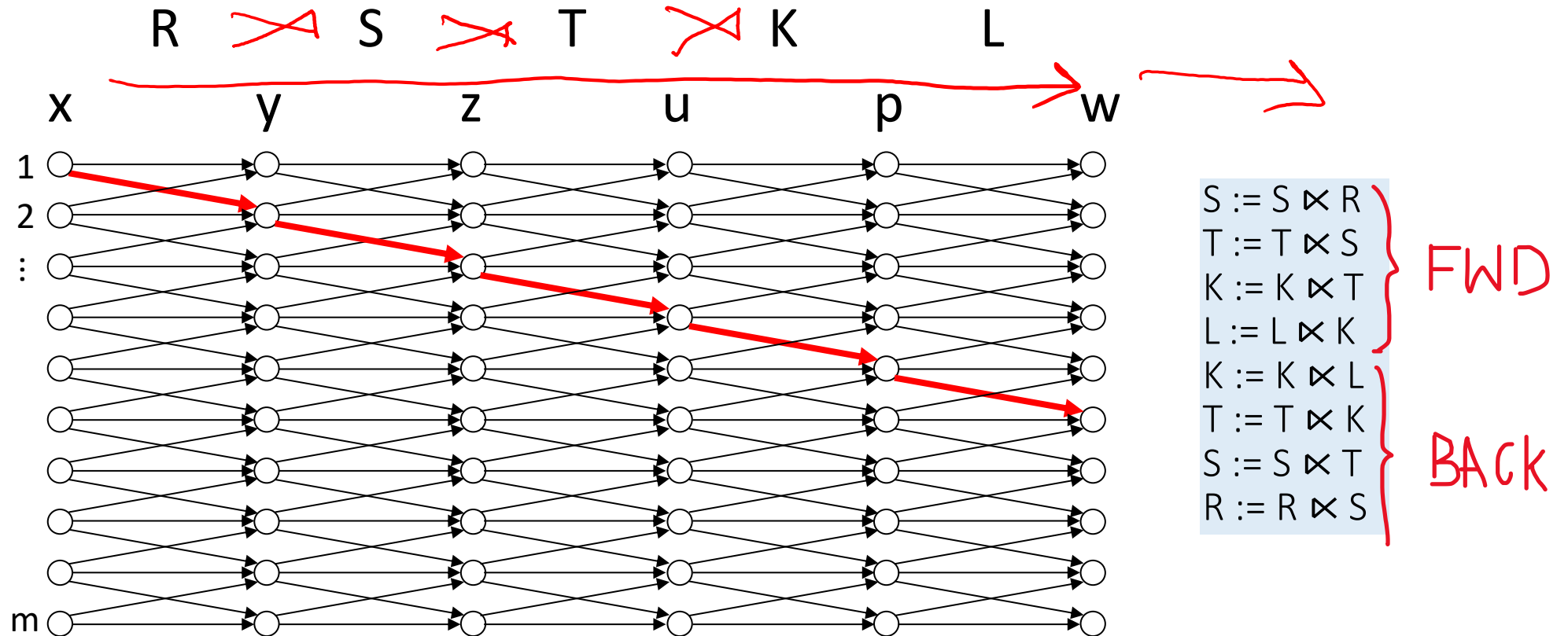
- Intermediate relations of size up to m^5 (m is here domain size!)
- But final answer can be as small as 0 (or 1...)

Application

$$\pi_x = \pi_x$$

$$Q_5^\infty(x,y,z,u,v,w) :- R(x,y), S(y,z), T(z,u), K(u,v), L(v,w).$$

Semi-join reduction allows pre-processing in $O(n)$ where n is size of DB (= number of edges)



- Intermediate relations of size up to m^5 (m is here domain size!)
- But final answer can be as small as 0 (or 1...)

What is so special about acyclic queries? Local = Global consistency

- "Consistency" = no dangling tuples (recall semi-join)

- **locally join consistent** (every tuple participates in pairwise joins)

$$\pi_{R_i}(R_i \bowtie R_j) = R_i$$

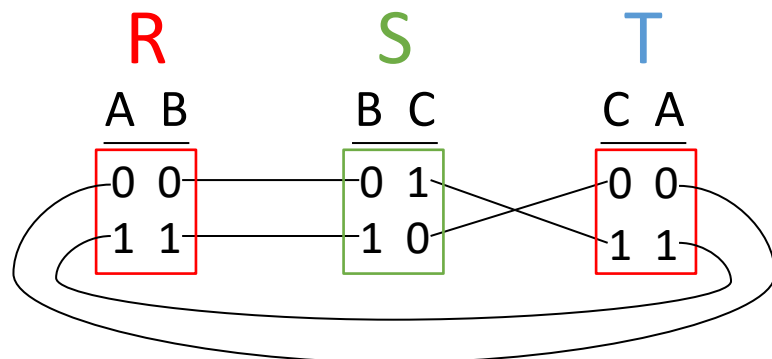
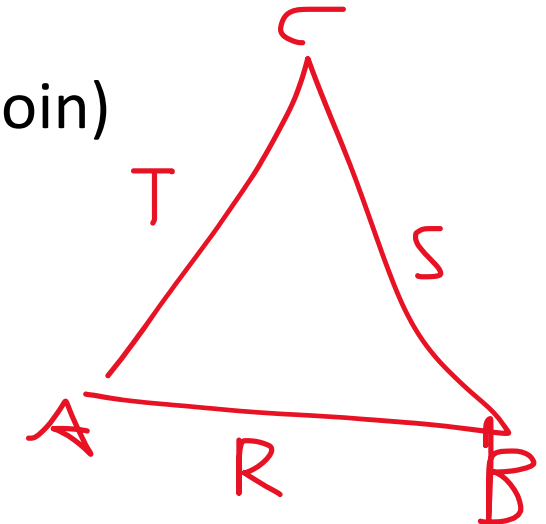
- **globally join consistent** (every tuple participates in the full join)

$$\pi_{R_i}(R_1 \bowtie R_2 \bowtie \dots \bowtie R_k) = R_i$$

- Both imply each other only for acyclic queries

- For all queries: global \Rightarrow local

- Acyclic queries: **local \Rightarrow global**; but not for cyclic queries!

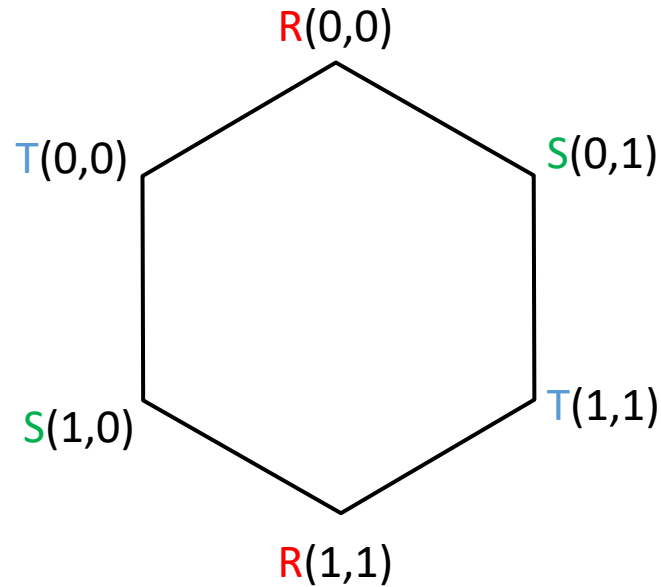


$$Q_{\Delta}(A, B, C): \neg R(A, B), S(B, C), T(C, A)$$

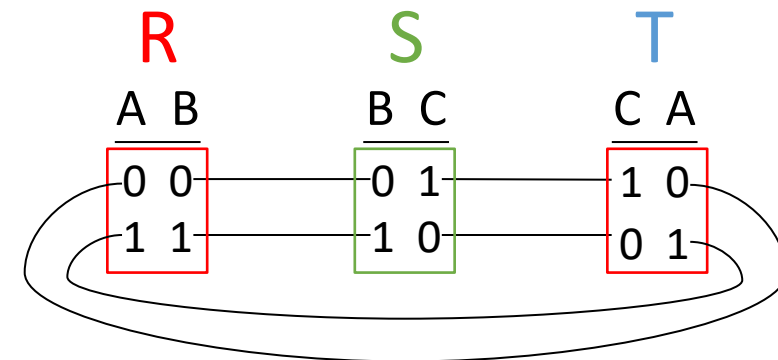
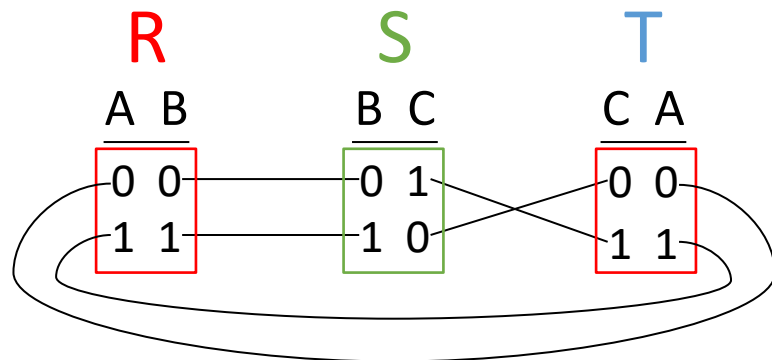
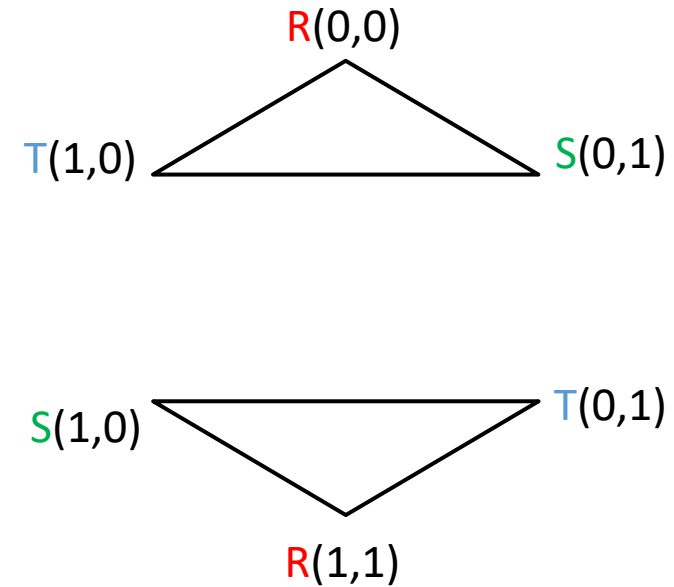
- Any two relations are locally consistent. E.g. $R \bowtie S$ is $\{(0,0,0), (1,1,1)\}$, which projected onto R is $\{(0,0), (1,1)\}$
- But $R \bowtie S \bowtie T = \emptyset$, so the relations are not globally consistent

What is so special about acyclic queries? Local = Global consistency

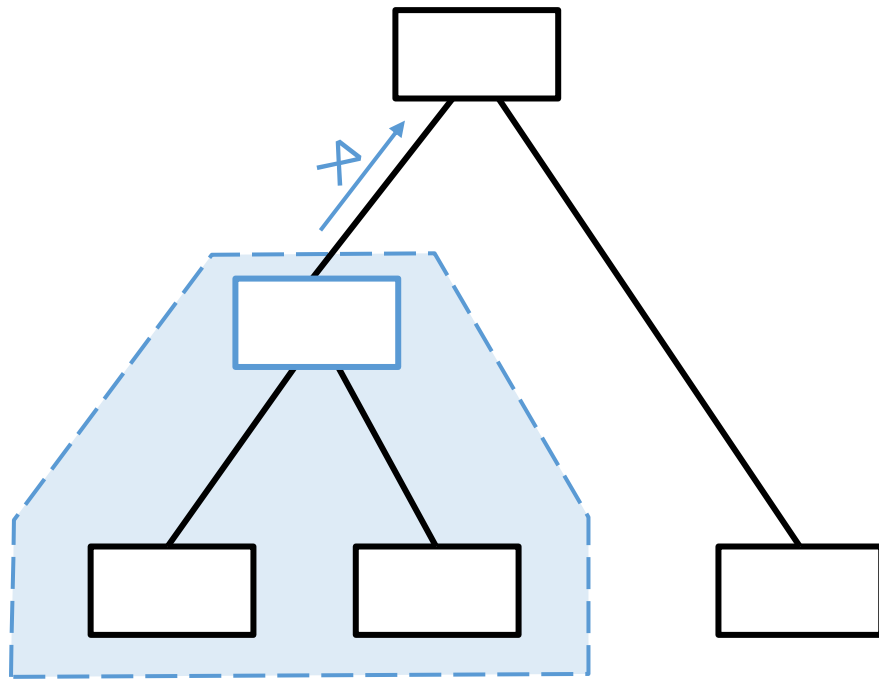
$$Q_{\Delta}(A, B, C): \neg R(A, B), S(B, C), T(C, A)$$



Semi-joins can't distinguish between the two examples. But only the right one is globally consistent



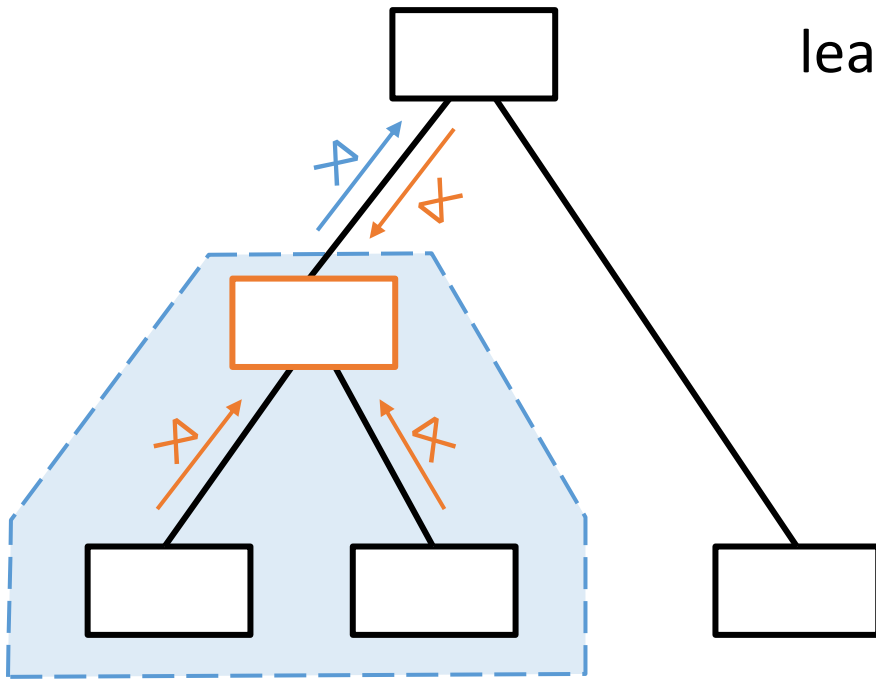
Semijoin Reduction as Message Passing on Trees



1. A message sent across an edge **contains all the information** from the subtree rooted at the sender.
⇒ Thus the reduction always needs to start at the leaves!
2. Key for acyclic queries: When joining, if there are no dangling tuples (Thus for a "reduced database") **every additional join can never decrease the size** of the intermediate query results!

Semijoin Reduction as Message Passing on Trees

Every **relation (node)** will be reduced d times where d is the degree of the node. Thus leaves (also the root) are reduced only once)



Each edge transmits **2 messages**: one up and one down.

Topic 3: Efficient query evaluation

Unit 1: Acyclic query evaluation

Lecture 21

Wolfgang Gatterbauer

CS7240 Principles of scalable data management (sp24)

<https://northeastern-datalab.github.io/cs7240/sp24/>

4/2/2024

Pre-class conversations

- Last class summary
- Project: Feedback by tomorrow WED
 - we can iterate, just approach me
- Scribes: Feedback by next week WED
 - please see my detailed comments, do also approach me after class with comments / questions / pointers. The goal is for the scribes and comments to be starters

- Today:
 - Yannakakis is instance optimal, but we do even better

Preface to "Theories of computability" by Pippenger (1997)

The central theme of this book is computability, in that most of the questions studied concern whether or not it is possible to perform some computation by some particular means (rather than how hard or easy it might be to do so).

Two kinds of results can be identified in almost any mathematical discipline. They do not seem to have standardized names, and so I shall propose the names “structural” and “cautionary.” It would be hopeless to attempt precise definitions, and so I shall try to epitomize the distinction with an example: the unique factorization of integers is a structural result, whereas the existence of number rings in which unique factorization fails is a cautionary result. Structural results show that the world (or at least the part being studied) is neater than one might expect, by exhibiting hidden simplicities. Most of the stones in the great structural edifices of mathematics fall into this category: structure theorems, normal forms, classifications, and so forth. Cautionary results show that the world is more complex than one might expect; it is here that the pathologies of examples and counterexamples are found. These results offer guidance to those who seek structural results, warning them away from impossible quests, but they also offer occasions to celebrate the richness and diversity of mathematical phenomena.

Semi-join reducers fail with the triangle



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges

Join tree

?

Query hypergraph

?

Semi-join reducers fail with the triangle



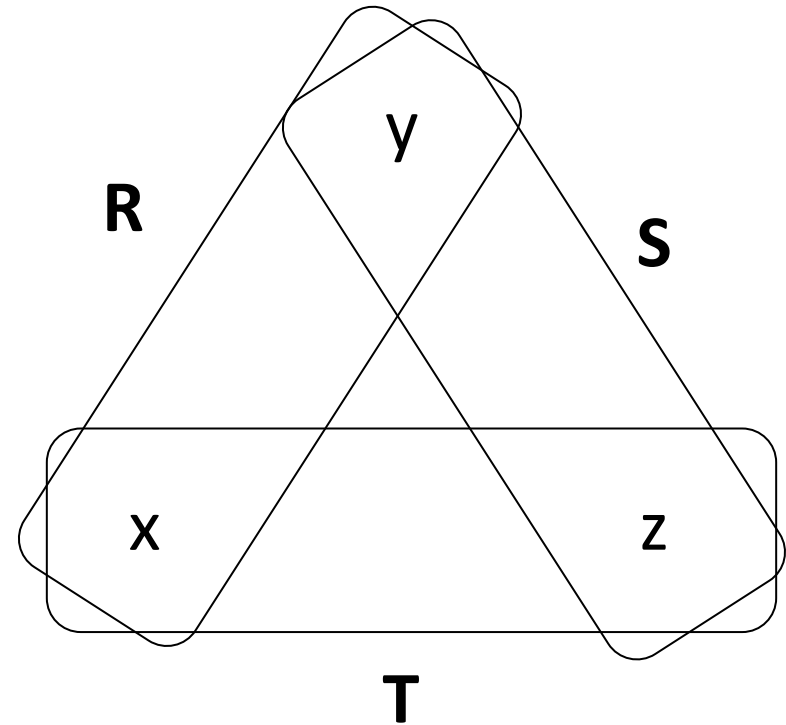
$Q(x,y,z) :- R(x,y), S(y,z), T(x,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges

Join tree

?



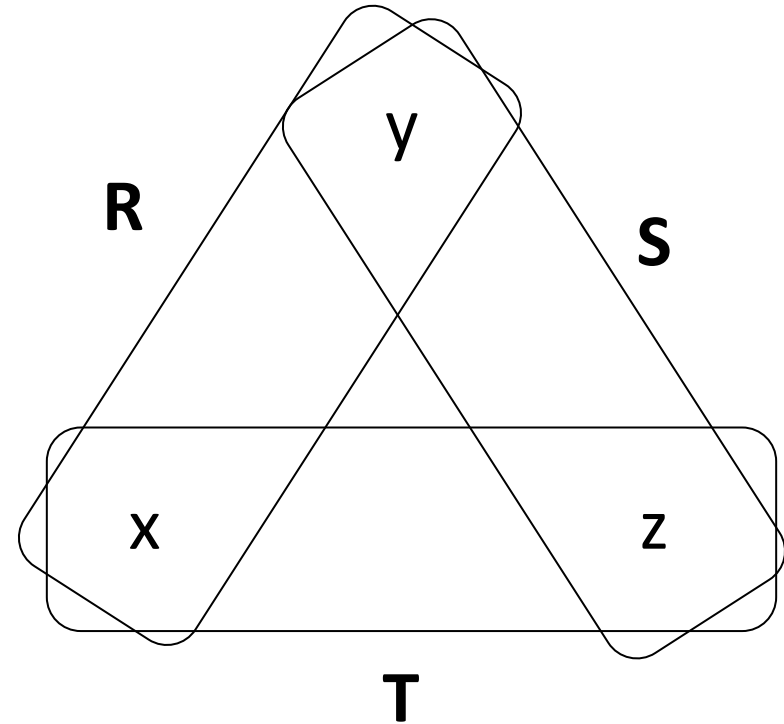
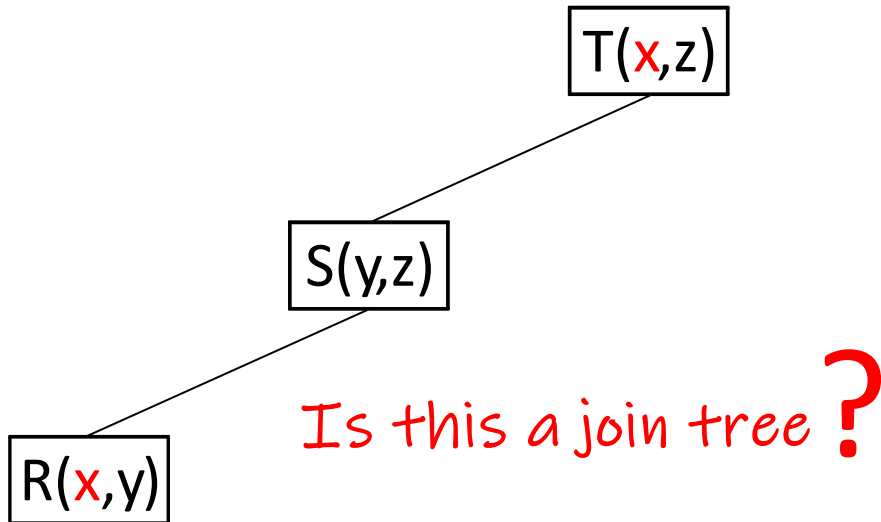
Semi-join reducers fail with the triangle



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges



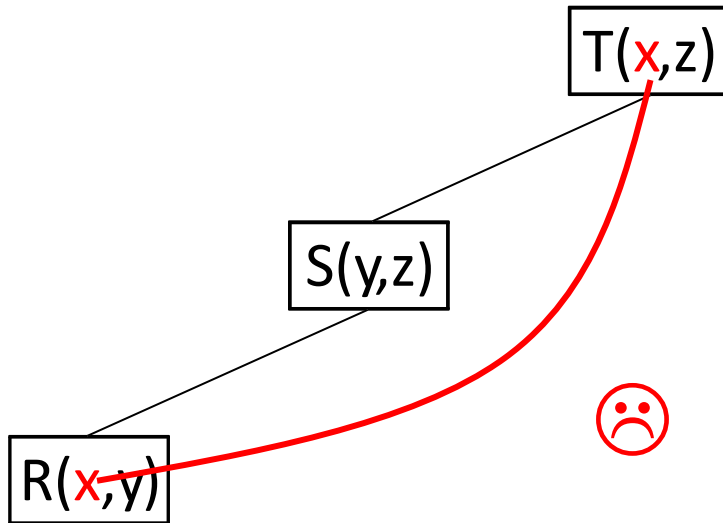
Semi-join reducers fail with the triangle



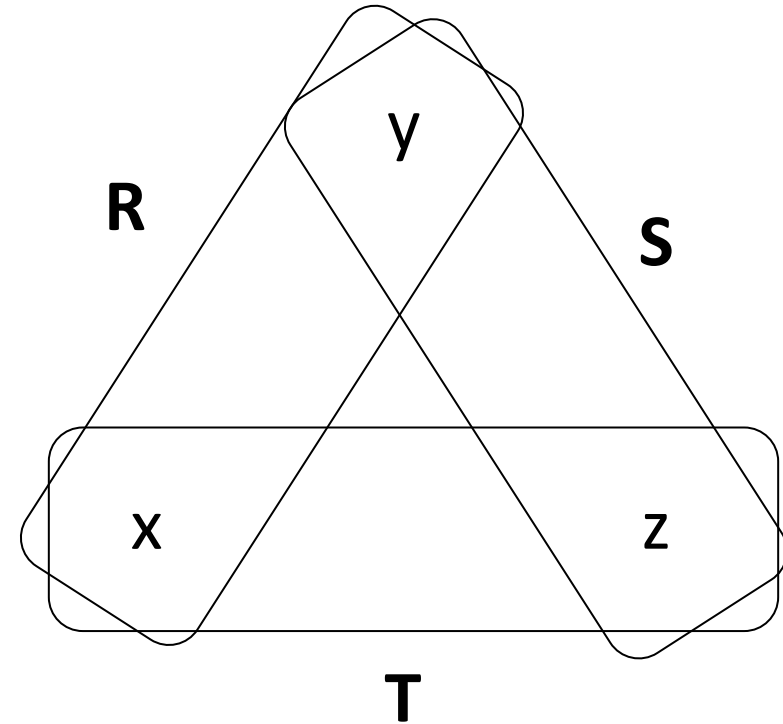
$Q(x,y,z) :- R(x,y), S(y,z), T(x,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges



There is no join tree! You can't fulfill the running intersection property...



Semi-join reducers work with the "beta-triangle"



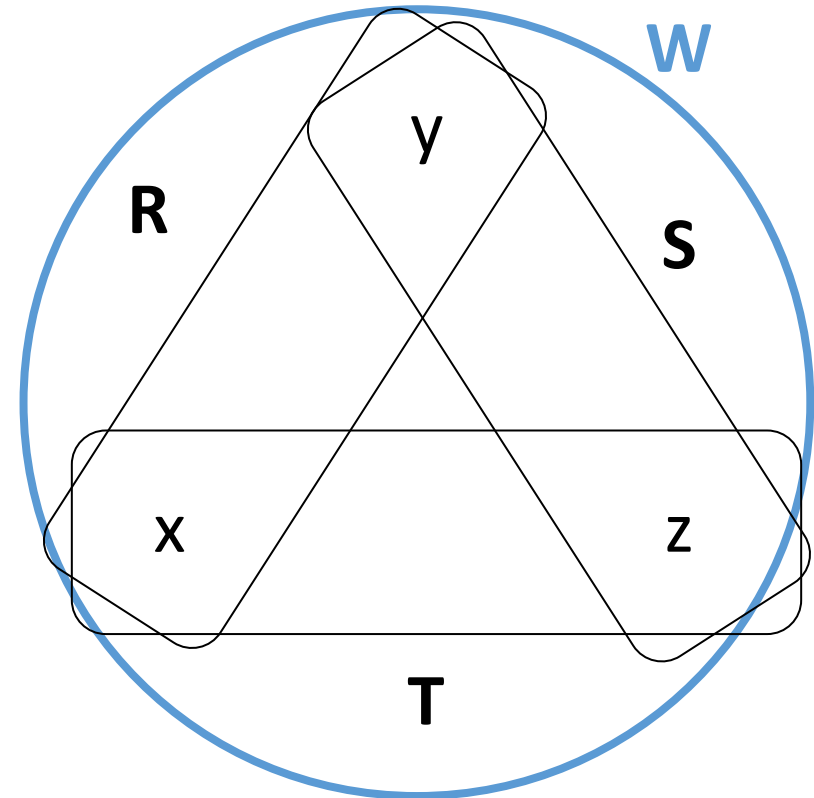
$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges

Join tree

?



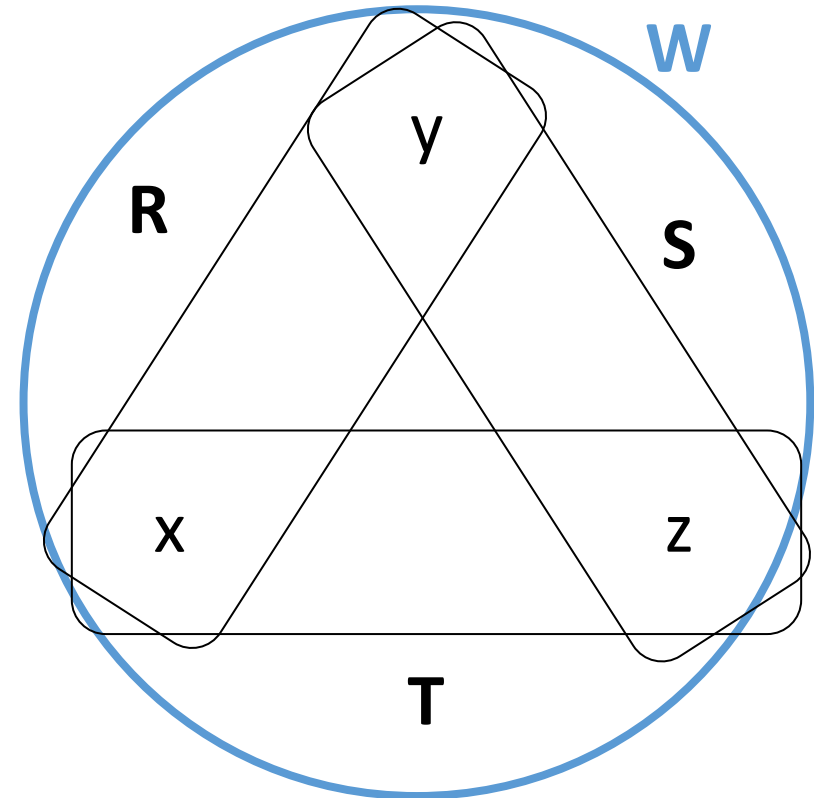
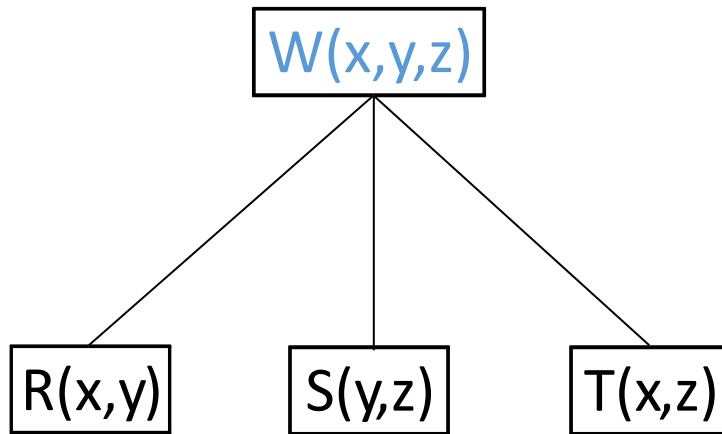
Semi-join reducers work with the "beta-triangle"



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$

GYO reduction (**ear removal**)

- remove isolated nodes
- remove consumed or empty edges



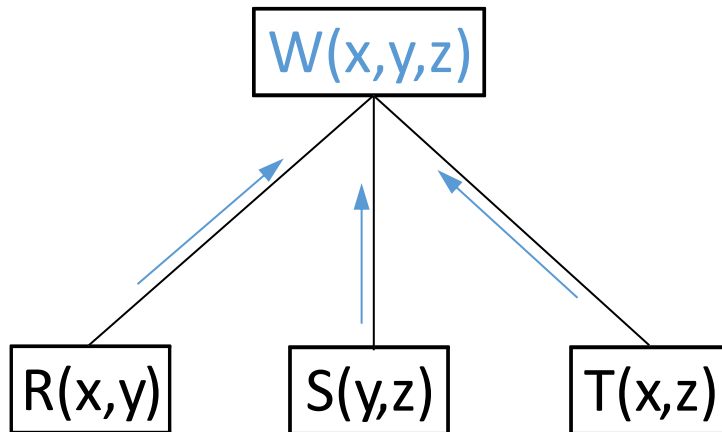
Next: Write 1) a full reducer and then 2) the new join expression in RA



Semi-join reducers work with the "beta-triangle"



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$



$$W_1(x, y, z) = W(x, y, z) \bowtie R(x, y)$$

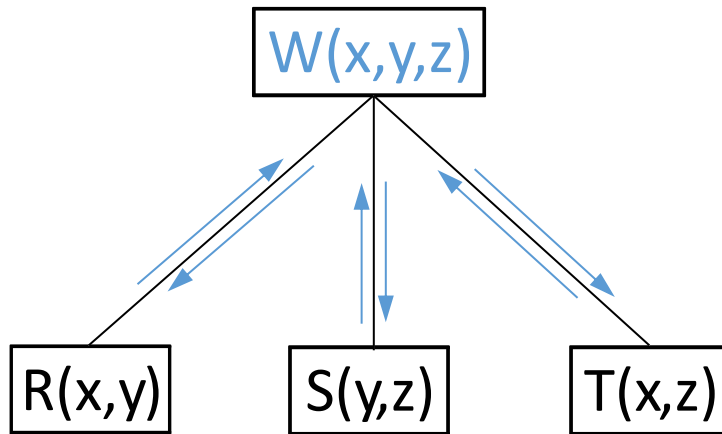
$$W_2(x, y, z) = W_1(x, y, z) \bowtie S(y, z)$$

$$W_3(x, y, z) = W_2(x, y, z) \bowtie T(x, z)$$

Semi-join reducers work with the "beta-triangle"



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$



$$W_1(x, y, z) = W(x, y, z) \bowtie R(x, y)$$

$$W_2(x, y, z) = W_1(x, y, z) \bowtie S(y, z)$$

$$W_3(x, y, z) = W_2(x, y, z) \bowtie T(x, z)$$

$$R_1(x, y) = R(x, y) \bowtie W_3(x, y, z)$$

$$S_1(y, z) = S(y, z) \bowtie W_3(x, y, z)$$

$$T_1(x, z) = T(x, z) \bowtie W_3(x, y, z)$$

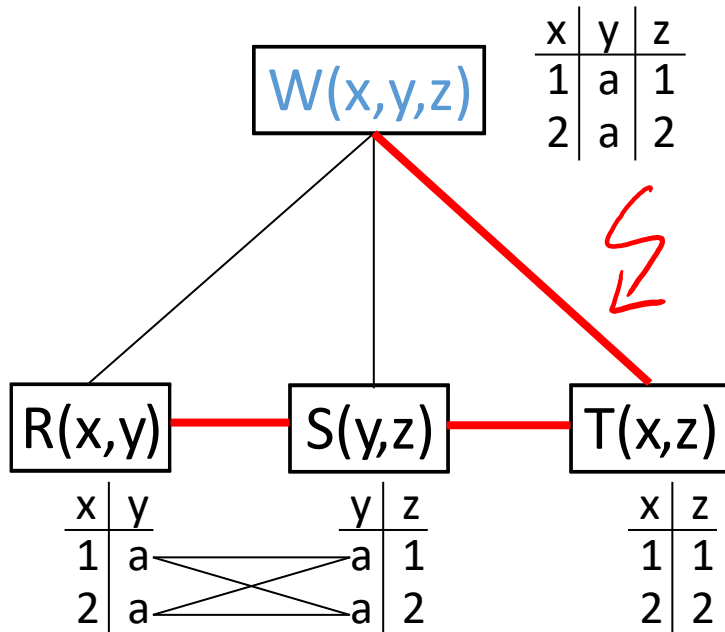
$Q(x,y,z) :- R_1(x,y), S_1(y,z), T_1(x,z), W_3(x,y,z).$

What can still go wrong after the reduction?

... but we still need to follow the join tree!



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$



$$\begin{aligned}
 W_1(x, y, z) &= W(x, y, z) \bowtie R(x, y) \\
 W_2(x, y, z) &= W_1(x, y, z) \bowtie S(y, z) \\
 W_3(x, y, z) &= W_2(x, y, z) \bowtie T(x, z) \\
 R_1(x, y) &= R(x, y) \bowtie W_3(x, y, z) \\
 S_1(y, z) &= S(y, z) \bowtie W_3(x, y, z) \\
 T_1(x, z) &= T(x, z) \bowtie W_3(x, y, z)
 \end{aligned}$$

$$Q(x,y,z) = \left(\left(R_1(x, y) \bowtie S_1(y, z) \right) \bowtie T_1(x, z) \right) \bowtie W_3(x, y, z)$$

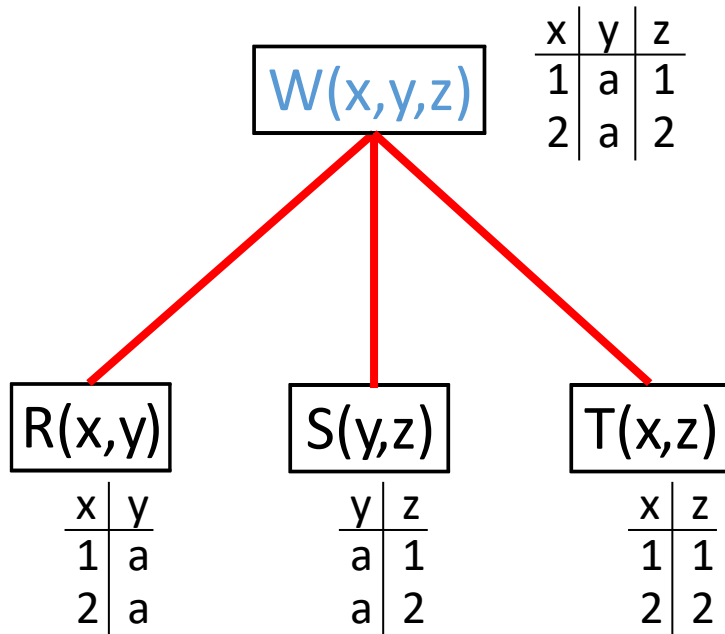
1	a	1
2	a	2
1	a	1
2	a	2

☹️ #=4

... but we still need to follow the join tree!



$Q(x,y,z) :- R(x,y), S(y,z), T(x,z), W(x,y,z).$



$$\begin{aligned}W_1(x, y, z) &= W(x, y, z) \bowtie R(x, y) \\W_2(x, y, z) &= W_1(x, y, z) \bowtie S(y, z) \\W_3(x, y, z) &= W_2(x, y, z) \bowtie T(x, z) \\R_1(x, y) &= R(x, y) \bowtie W_3(x, y, z) \\S_1(y, z) &= S(y, z) \bowtie W_3(x, y, z) \\T_1(x, z) &= T(x, z) \bowtie W_3(x, y, z)\end{aligned}$$

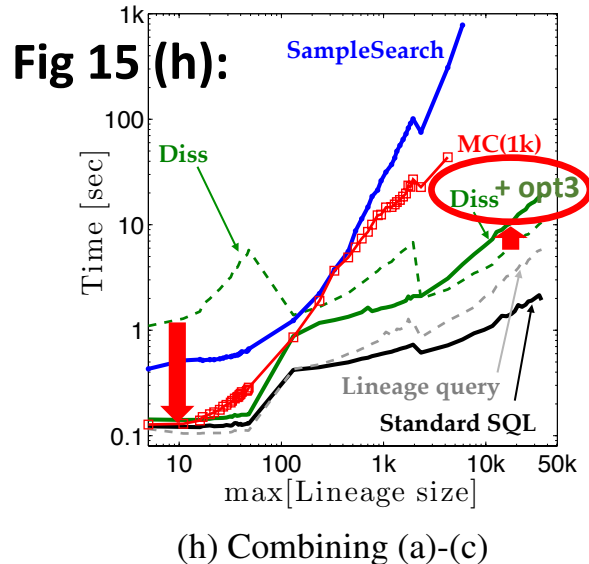
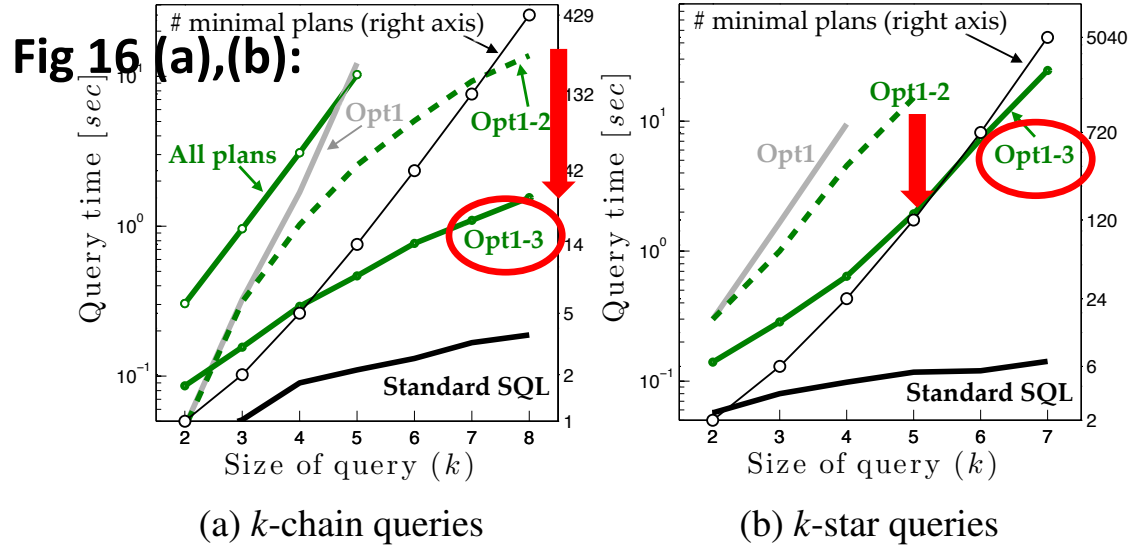
~~$Q(x,y,z) = ((R_1(x,y) \bowtie S_1(y,z)) \bowtie T_1(x,z)) \bowtie W_3(x,y,z)$~~

We still need to follow the join tree!



$$Q(x,y,z) = ((R_1(x,y) \bowtie W_3(x,y,z)) \bowtie S_1(y,z)) \bowtie T_1(x,z)$$

Semi-join reductions can be extremely powerful



Semi-join reductions can be extremely powerful in different contexts (yet is speculative, i.e. depend on the concrete input to pay off)

6.3 Opt. 3: Deterministic semi-join reduction

The most expensive operations in probabilistic query plans are the group-bys for the probabilistic project operations. These are often applied early in the plans to tuples which are later pruned and do not contribute to the final query result. Our third optimization is to first apply a *full semi-join reduction on the input relations* before starting the probabilistic evaluation from these *reduced input relations*.

We like to draw here an important connection to [54], which introduces the idea of “lazy plans” and shows orders of magnitude performance improvements for safe plans by computing confidences not after each join and projection, but rather at the very end of the plan. We note that our semi-join reduction *serves the same purpose* with similar performance improvements and also apply for safe queries. The advantage of semi-join reductions, however, is that we *do not require any modifications to the query engine*.

Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

<https://northeastern-datalab.github.io/>

Key idea of [Yannakakis'81]: Reduction by Semi-joins

- Key insight for acyclic queries:
 - If there are no dangling tuples, then the result can never shrink with an additional join
 - Thus, for a "reduced database", every additional join can only increase the size of the intermediate query results

Yannakakis Algorithm [Yannakakis'81]

- Given: acyclic full conjunctive query Q (full = no projections)
- Compute Q on any database in time $O(|\text{Input}| + |\text{Output}|)$ by using any rooted **join tree**
- Step 1: **semi-join reduction** (two sweeps)
 - Pick any root node R in the join tree of Q
 - Step 1a: Do a semi-join reduction from the leaves to R (bottom-up)
 - Step 1b: Do a semi-join reduction from R to the leaves (top-down)
- Step 2: use the **join tree as query plan**: pick any root and join bottom-up or top-down
 - Notice that step 2 can be combined with the top-down SJ-reduction

[Yannakakis'81] Acyclic Conjunctive Queries

- Theorem [Yannakakis'81]:
 - The Acyclic Conjunctive Query Evaluation Problem is tractable.
 - There is an algorithm for query evaluation with following properties:
 - If Q is a Boolean acyclic conjunctive query, then the algorithm runs in time: $O(|Q| \cdot |D|)$
 - If Q is a full (i.e. no projections) acyclic conjunctive query then the algorithm runs in time: $O(|Q| \cdot |D| + |Q(D)|)$
 - In terms of data complexity, this means $O(|Input| + |Output|)$
 - i.e., it runs in **input/output linear time**, which is the “right” notion of tractability in this case (you can't do better, in general)

Yannakakis' Algorithm for Boolean acyclic CQs

Basically the standard **Dynamic Programming** Algorithm

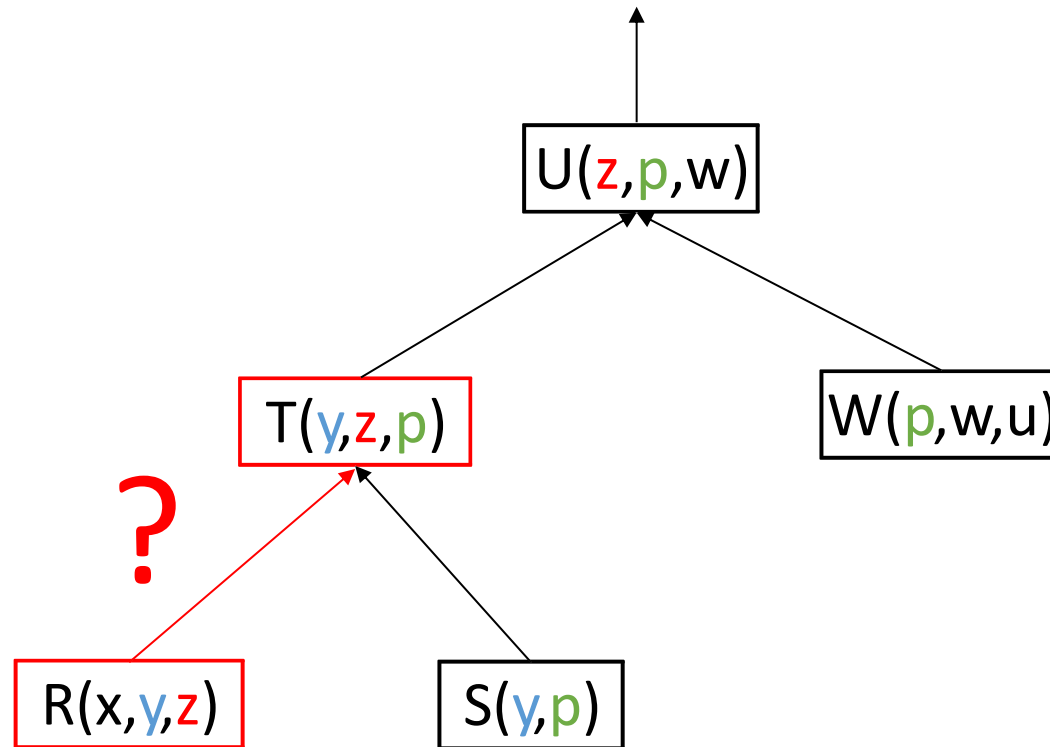
- Input: Boolean acyclic conjunctive query Q , database D
- Construct a **join tree** T of Q
- Populate the nodes of T with the matching relations of D .
- Traverse the tree T bottom up (also called "collection phase"):
 - For each node compute the **semi-joins** of the (current) relation in the node with the (current) relations in the children of the node
- Examine the resulting relation R at the root of T
 - If R is non-empty, then output $Q(D) = 1$ (D satisfies Q).
 - If R is empty, then output $Q(D) = 0$ (D does not satisfy Q).

Acyclic Conjunctive Queries



- Where are the semi-join reductions in following query:

$Q :- R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$



Rooted Join Tree for Q

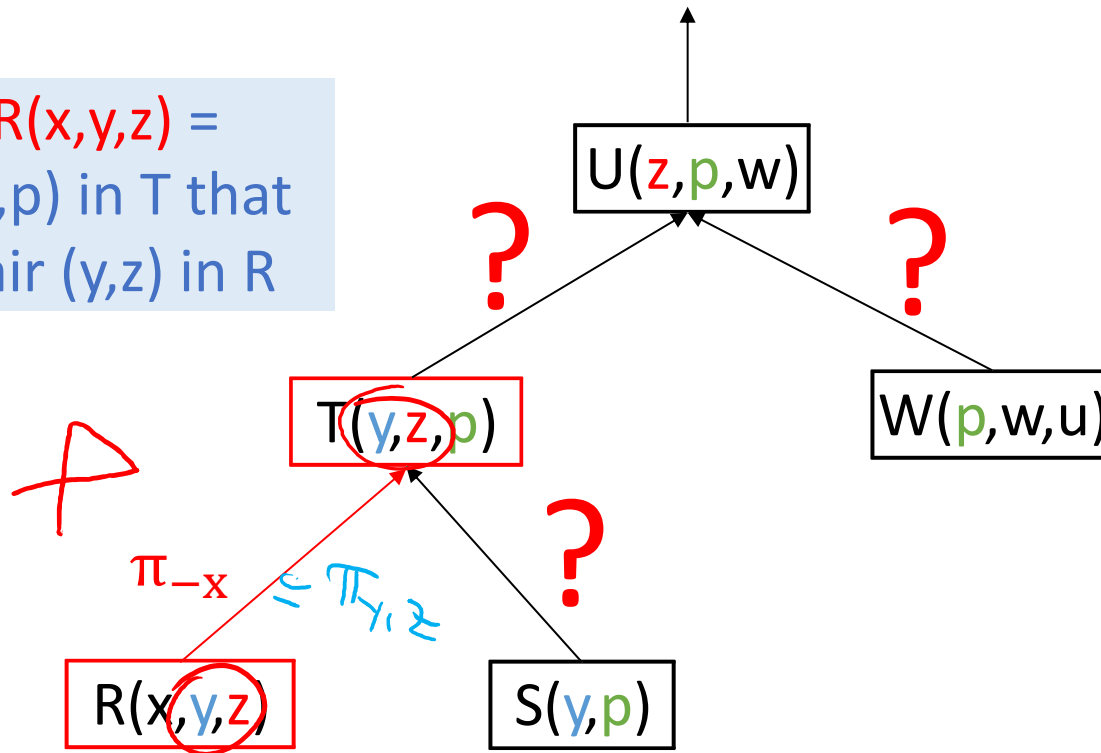
Acyclic Conjunctive Queries



- Where are the semi-join reductions in following query:

$Q \text{ :- } R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$

$T(y,z,p) \bowtie R(x,y,z) =$
all triples (y,z,p) in T that
"match" a pair (y,z) in R



Rooted Join Tree for Q

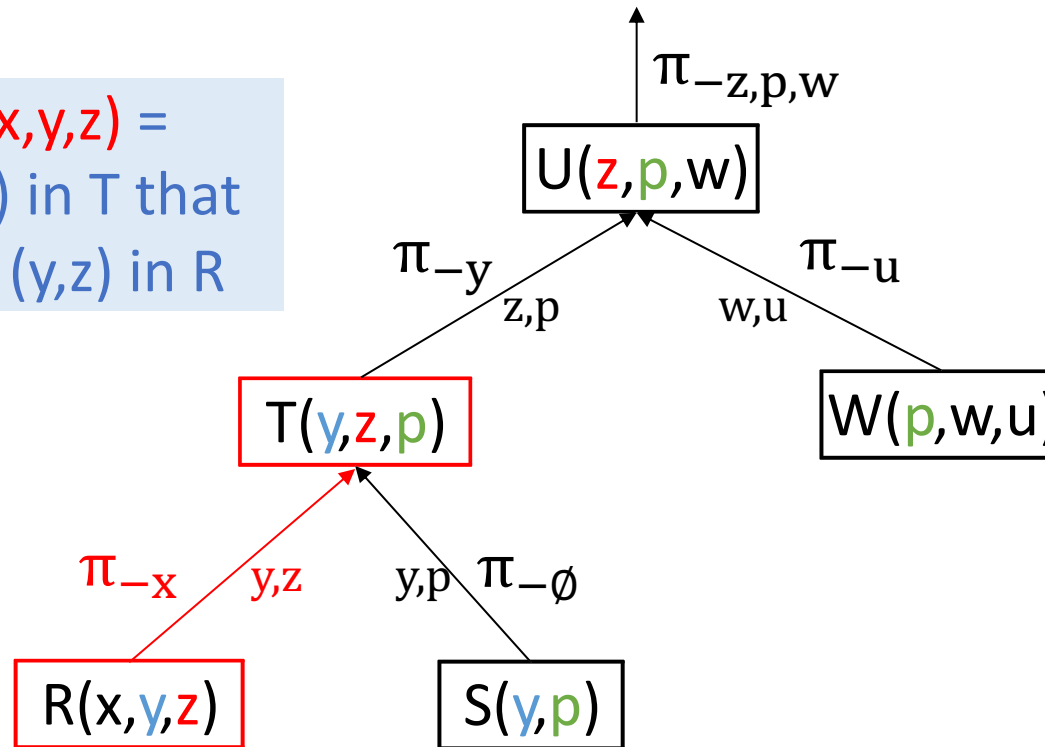
Acyclic Conjunctive Queries



- Where are the semi-join reductions in following query:

$Q \text{ :- } R(x,y,z), S(y,p), T(y,z,p), U(z,p,w), W(p,w,u).$

$T(y,z,p) \bowtie R(x,y,z) =$
all triples (y,z,p) in T that
"match" a pair (y,z) in R



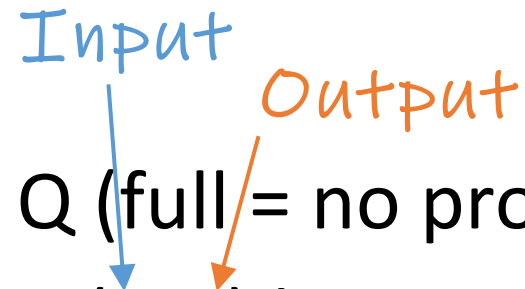
Rooted Join Tree for Q

More on Yannakakis' Algorithm

- The **join tree** makes it possible to avoid exponential explosion (in size $|Q|$) of the size of the intermediate computations.
- The algorithm can be extended to non-Boolean conjunctive queries using one additional top-down traversal of the join tree.
 - Bottom up (or "**collect**")
 - Top down (or "**distribute**")
- There are efficient algorithms for detecting acyclicity & computing a join tree.
 - [Tarjan, Yannakakis'84] Linear-time algorithm for detecting acyclicity and computing a join tree.
 - [Gottlob, Leone, Scarcello'01 (FOCS'98)] Detecting acyclicity is LOGCFL-complete and thus highly parallelizable)

A detailed example for
the Yannakakis algorithm

Yannakakis Algorithm



- Given: acyclic full conjunctive query Q (full = no projections)
- Compute Q on any database in time $O(n+r)$ by using the **join tree**
- Step 1: **semi-join reduction** (two sweeps)
 - Pick any root node R in the join tree of Q
 - Sweep **bottom-up**: Do a semi-join reduction from the leaves to R
 - Sweep **top-down**: Do a semi-join reduction from R to the leaves
- Step 2: use the **join tree as query plan**: pick any root and join bottom-up or top-down
 - Notice that step 2 can be combined with the top-down SJ-reduction

Yannakakis example: first use GYO to get the join tree

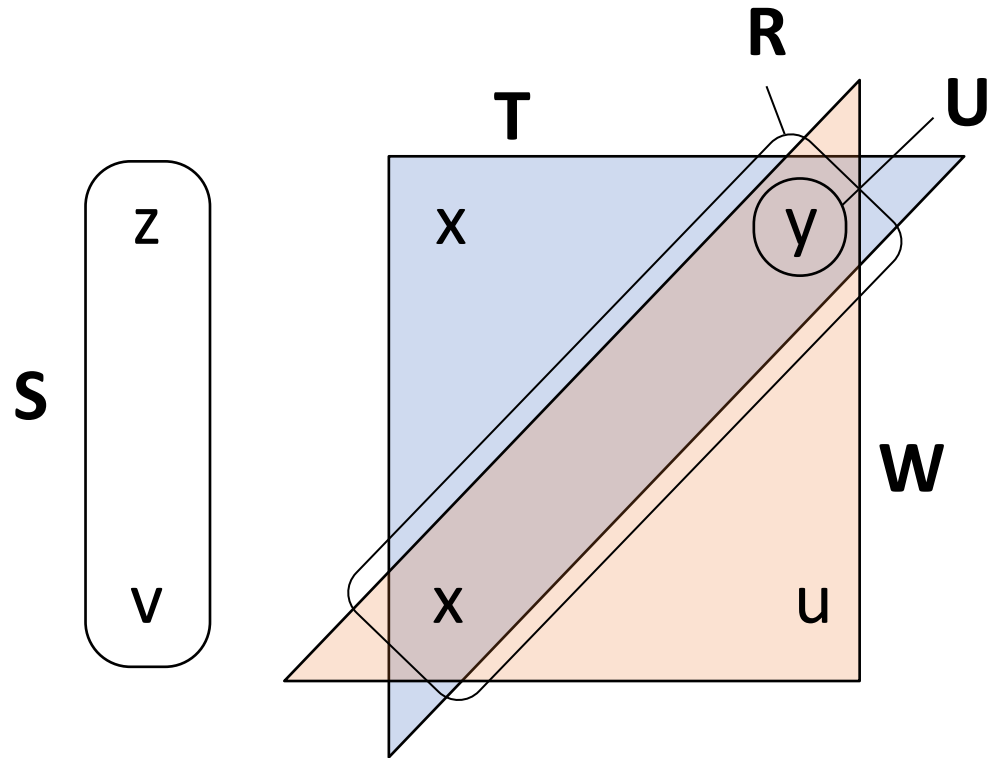
$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

R(x,y)	
x	y
a ₁	b ₁
a ₁	b ₂
a ₄	b ₆

one of several possible ones

Rooted Join tree

Hypergraph



S(z,v)	
z	v
c ₁	d ₁
c ₁	d ₂
c ₄	d ₆

T(p,x,y)		
p	x	y
e ₁	a ₁	b ₁
e ₁	a ₁	b ₂
e ₃	a ₃	b ₁
e ₃	a ₁	b ₄
e ₂	a ₂	b ₃

?

U(y)
y
b ₁
b ₂
b ₃

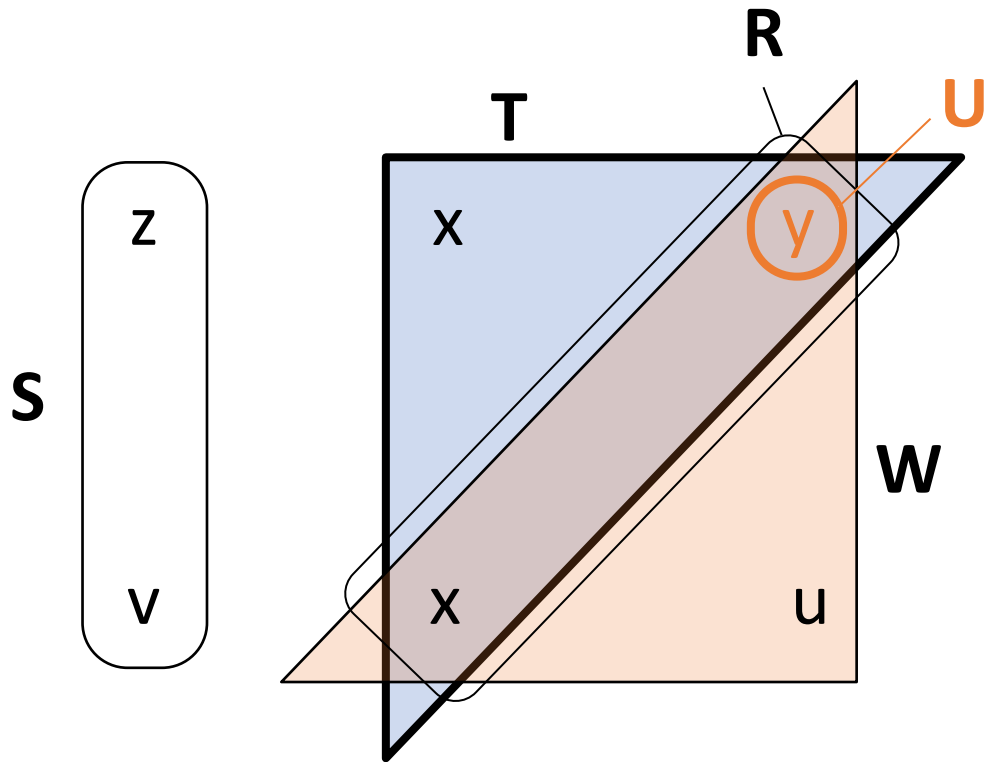
W(u,x,y)		
u	x	y
f ₁	a ₁	b ₁
f ₁	a ₁	b ₂
f ₂	a ₁	b ₂
f ₂	a ₂	b ₂

Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

R(x,y)	
x	y
a ₁	b ₁
a ₁	b ₂
a ₄	b ₆

Hypergraph



Rooted Join tree

S(z,v)	
z	v
c ₁	d ₁
c ₁	d ₂
c ₄	d ₆

T(p,x,y)		
p	x	y
e ₁	a ₁	b ₁
e ₁	a ₁	b ₂
e ₃	a ₃	b ₁
e ₃	a ₁	b ₄
e ₂	a ₂	b ₃

shared variable y
no isolated variable

U(y)
y
b ₁
b ₂
b ₃

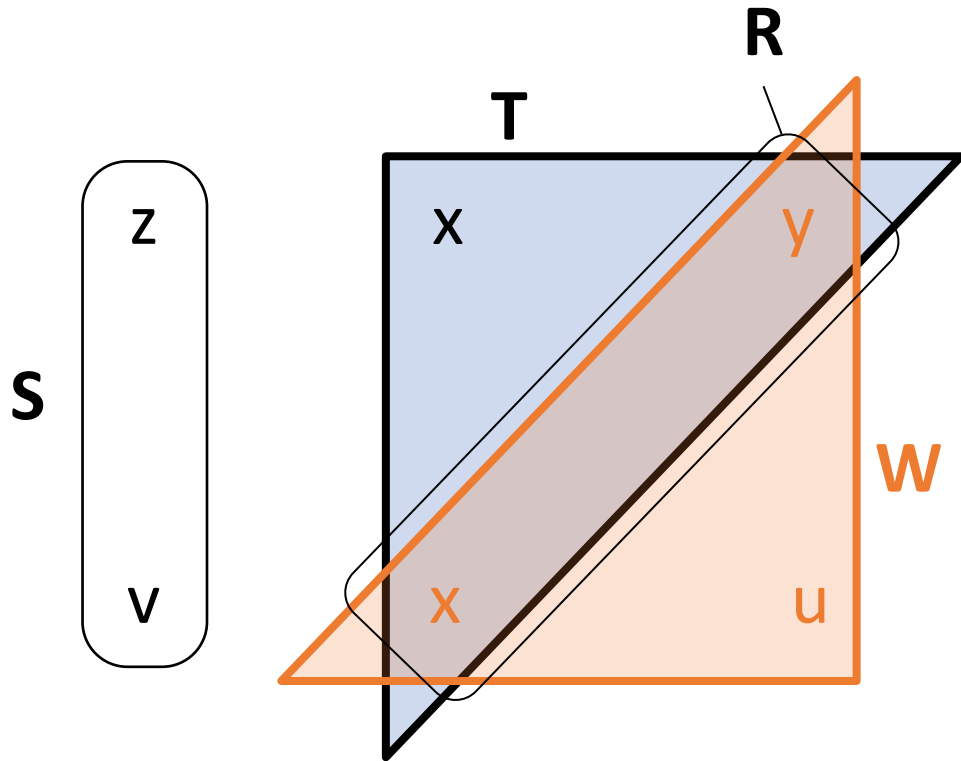
W(u,x,y)		
u	x	y
f ₁	a ₁	b ₁
f ₁	a ₁	b ₂
f ₂	a ₁	b ₂
f ₂	a ₂	b ₂

Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

R(x,y)	
x	y
a ₁	b ₁
a ₁	b ₂
a ₄	b ₆

Hypergraph



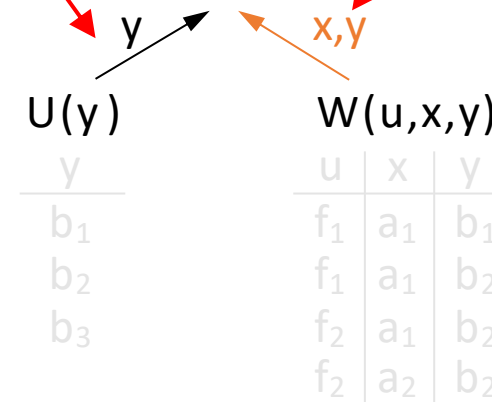
Rooted Join tree

S(z,v)	
z	v
c ₁	d ₁
c ₁	d ₂
c ₄	d ₆

T(p,x,y)		
p	x	y
e ₁	a ₁	b ₁
e ₁	a ₁	b ₂
e ₃	a ₃	b ₁
e ₃	a ₁	b ₄
e ₂	a ₂	b ₃

shared variable y
no isolated variable

shared variables x,y, isolated variable u gets projected away



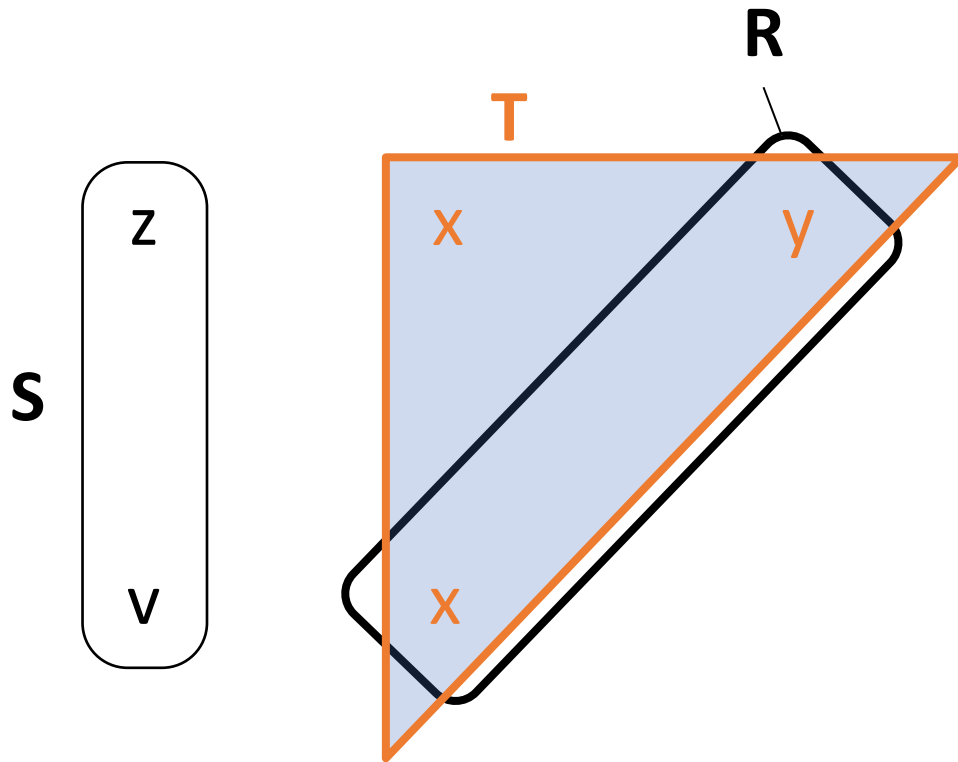
U(y)
y
b ₁
b ₂
b ₃

W(u,x,y)		
u	x	y
f ₁	a ₁	b ₁
f ₁	a ₁	b ₂
f ₂	a ₁	b ₂
f ₂	a ₂	b ₂

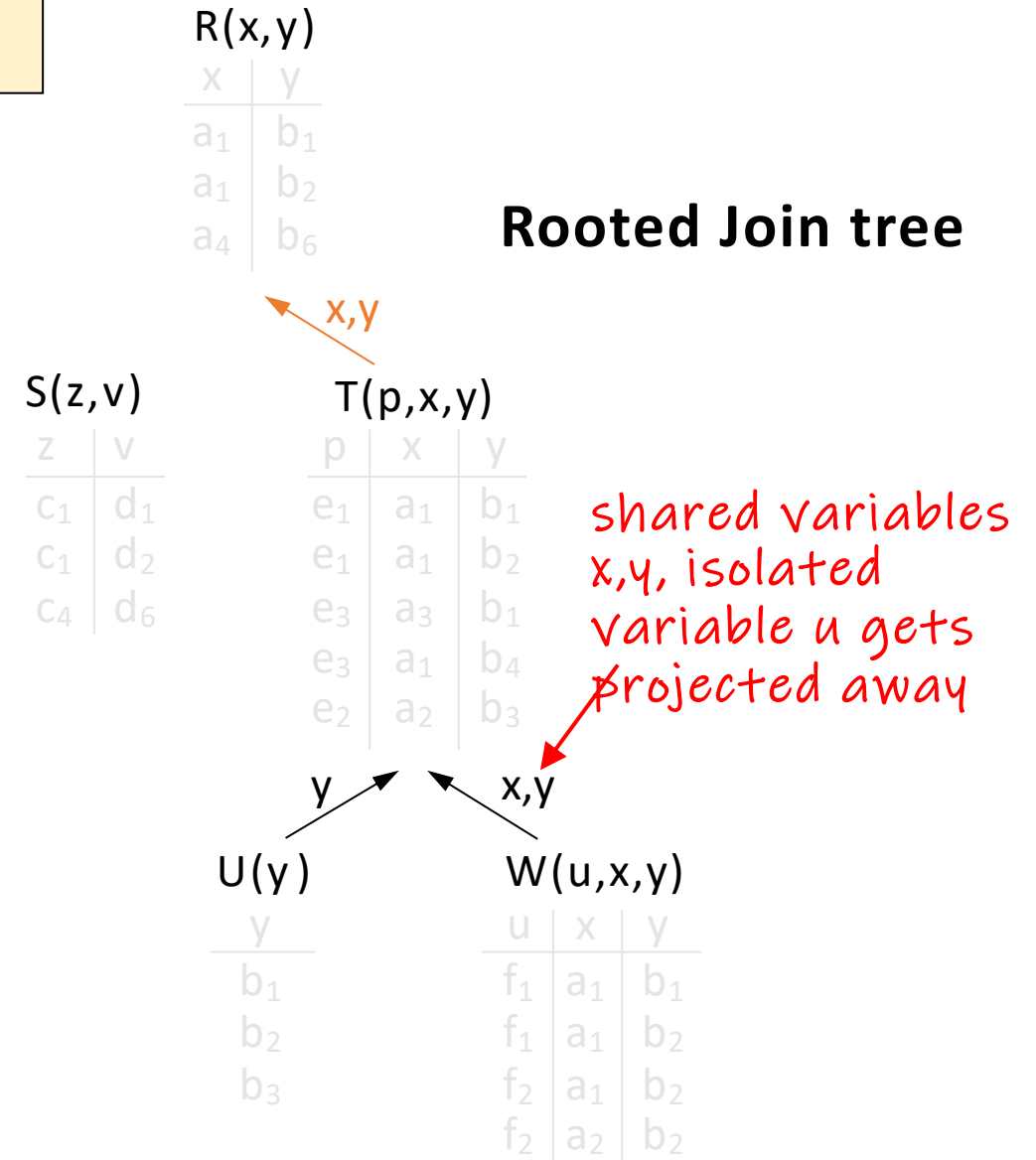
Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

Hypergraph



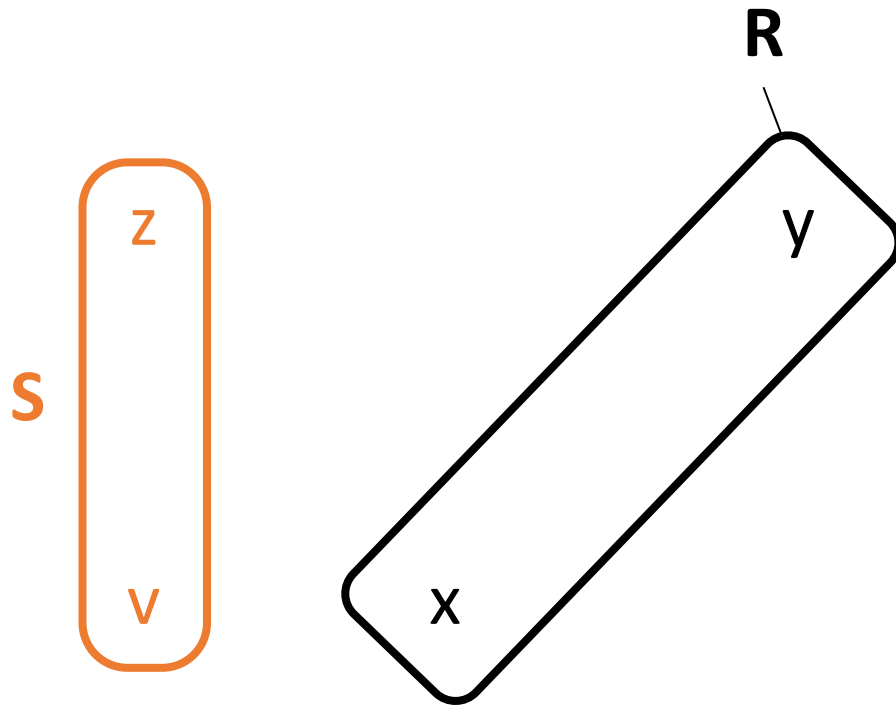
Rooted Join tree



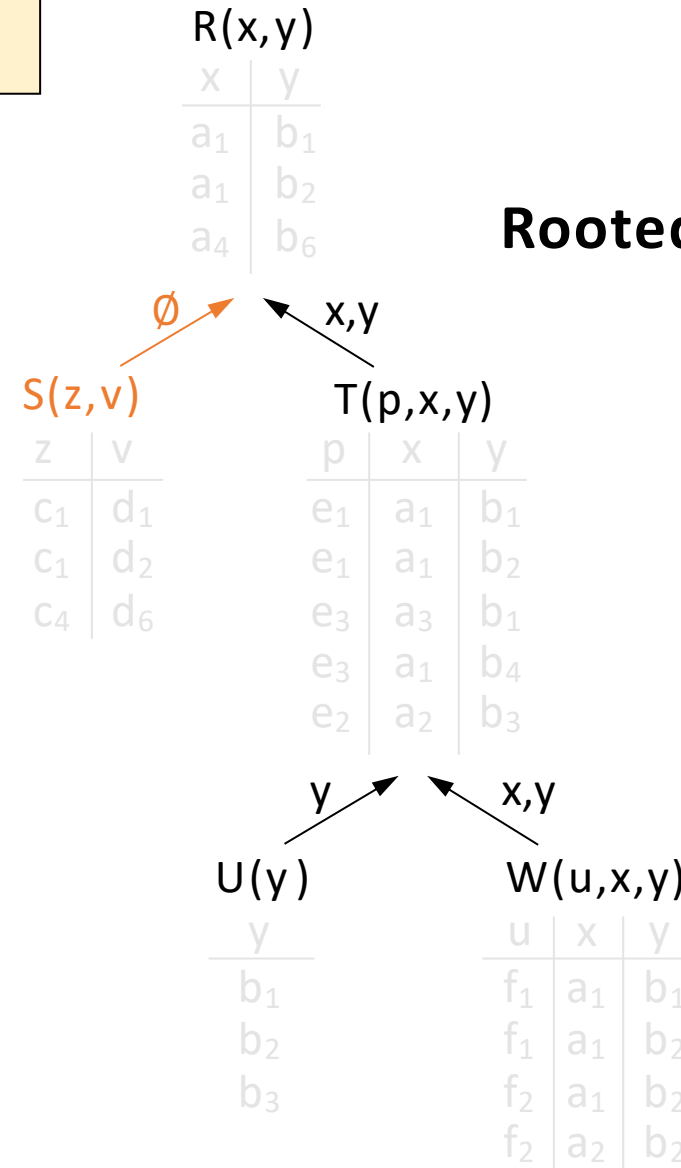
Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

Hypergraph



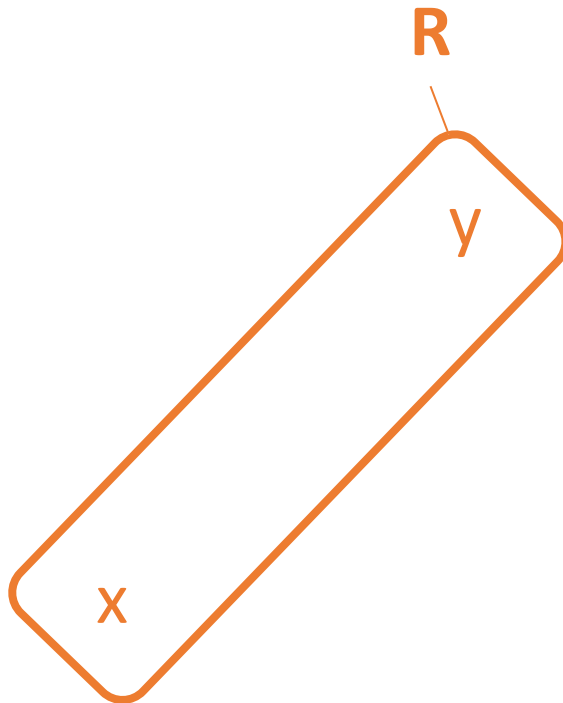
Rooted Join tree



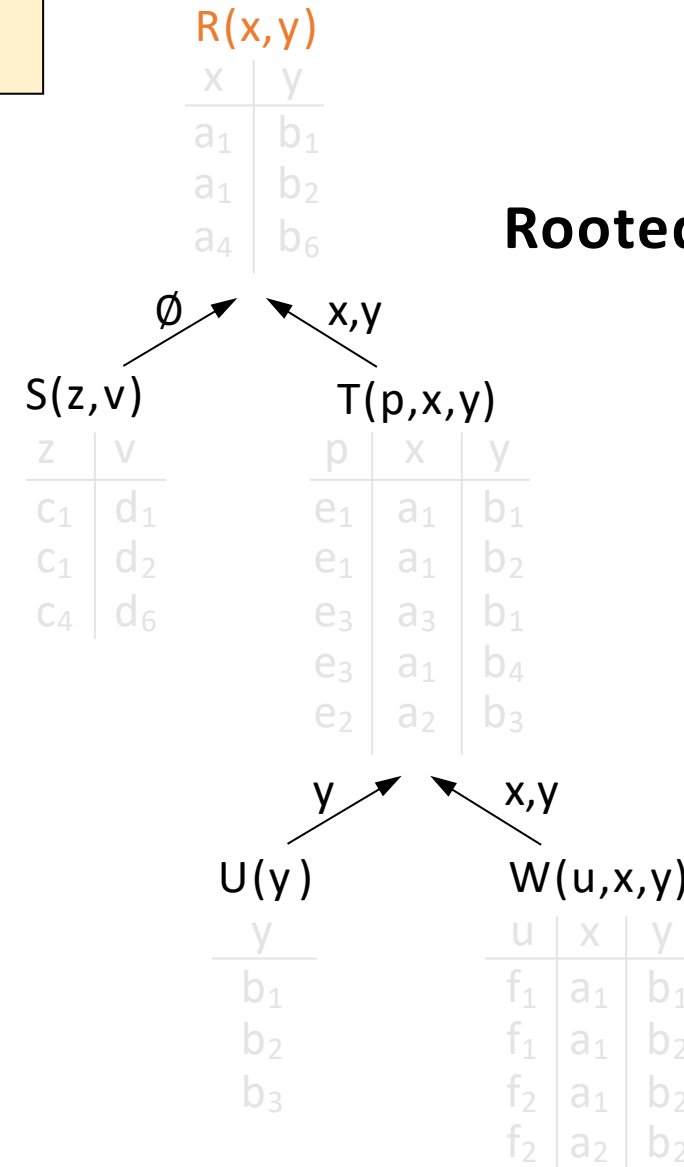
Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

Hypergraph

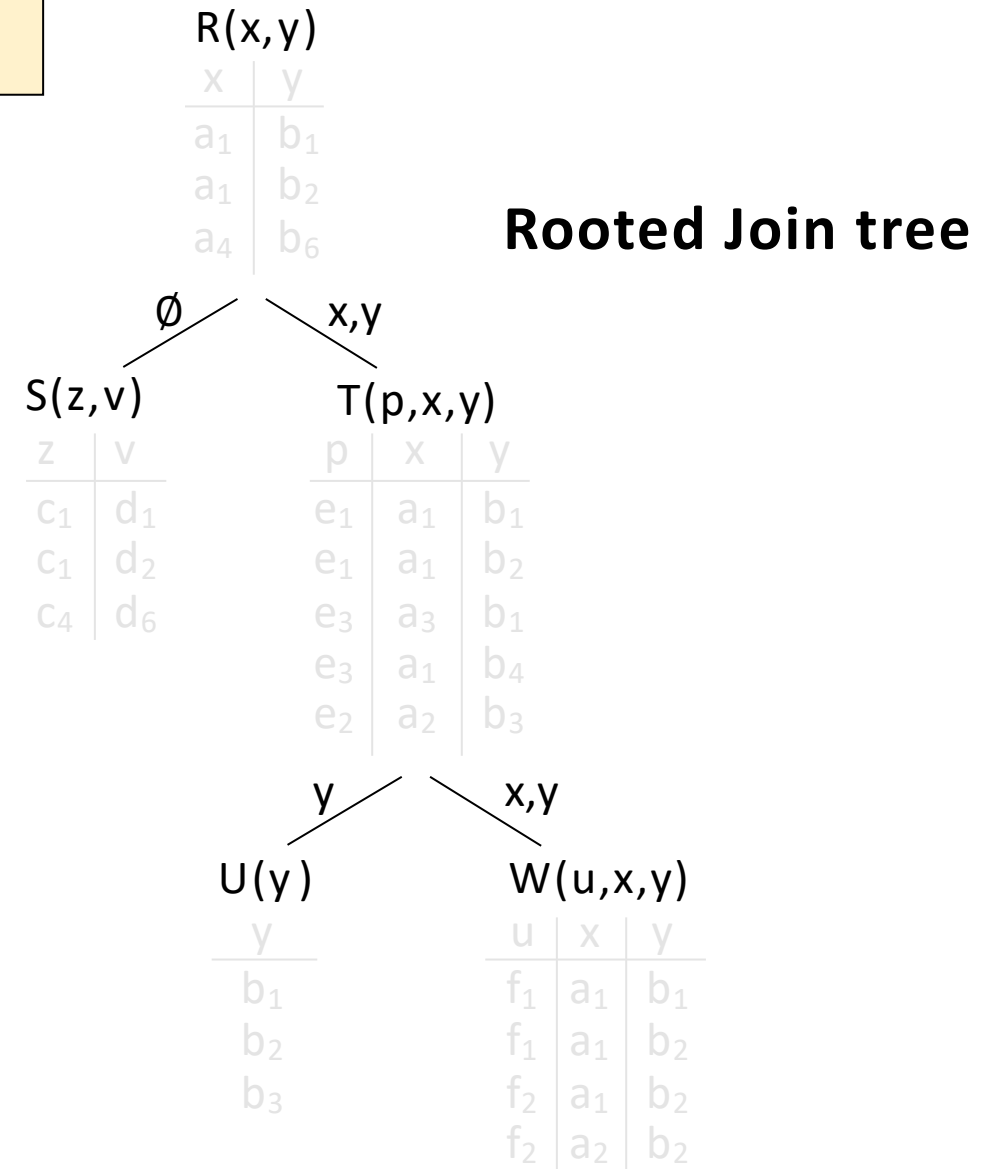


Rooted Join tree



Yannakakis example: first use GYO to get the join tree

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

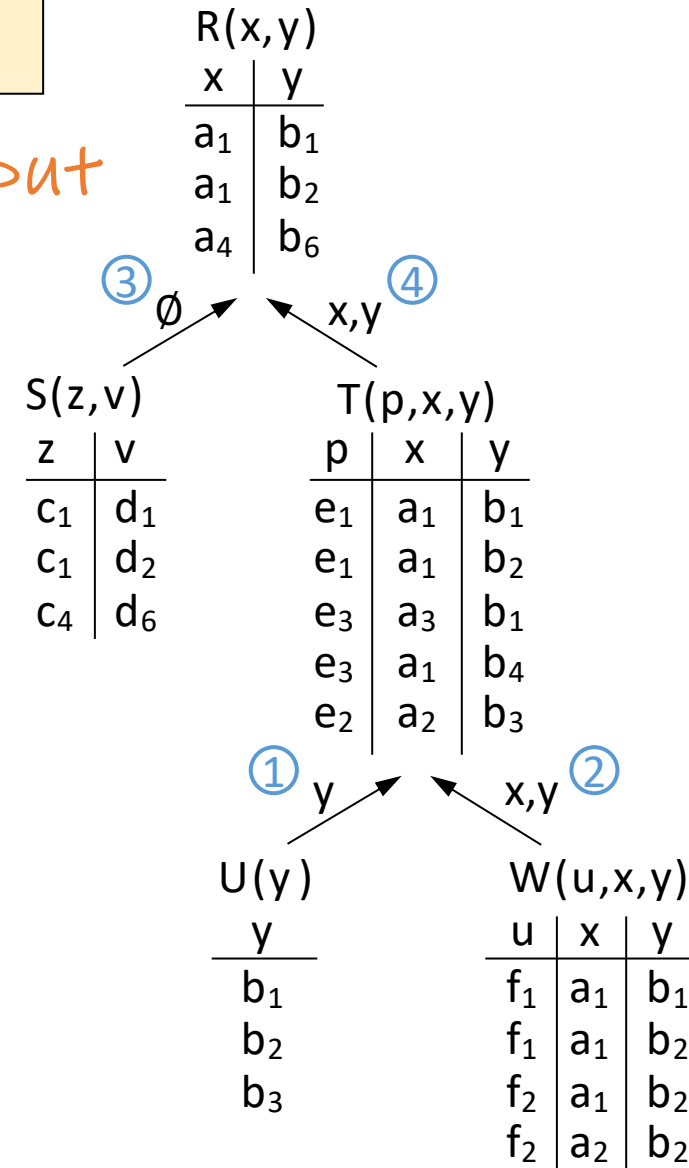


Yannakakis Algorithm example: 1st pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. **Semi-join phase** \times (remove dangling tuples) in $O(n)$ ← **Input**

- **Bottom-up semi-join propagation from leaves to root in some reverse topological order**

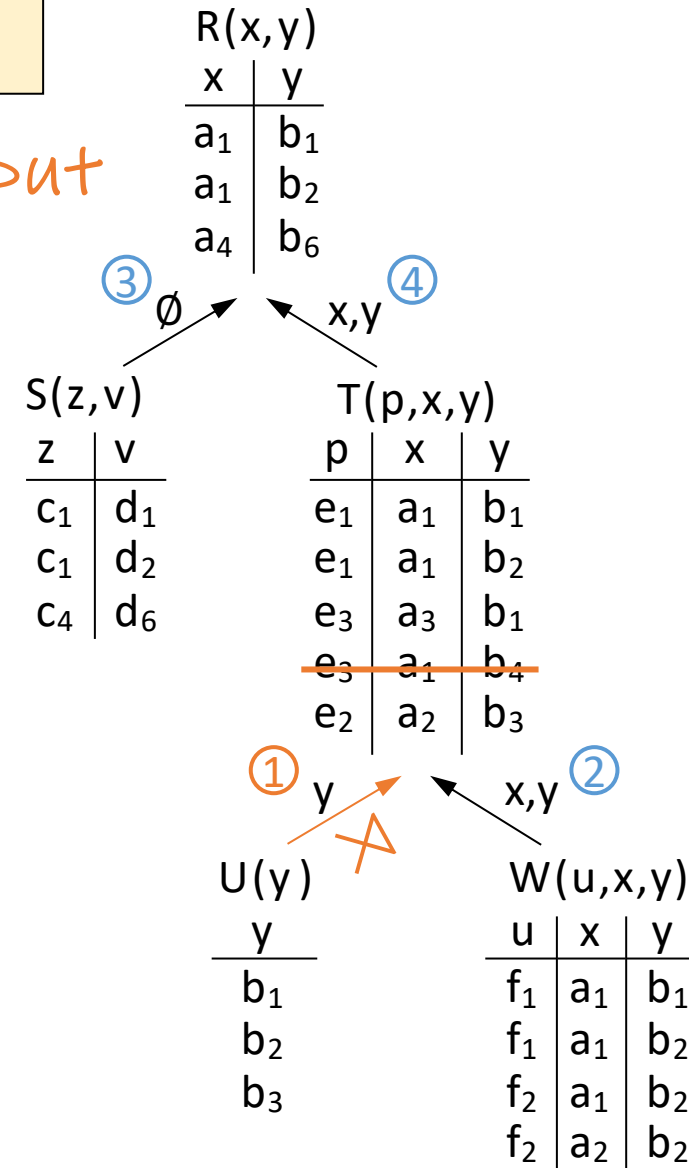


Yannakakis Algorithm example: 1st pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. **Semi-join phase** \times (remove dangling tuples) in $O(n) \leftarrow \text{Input}$

- **Bottom-up semi-join propagation from leaves to root in some reverse topological order**

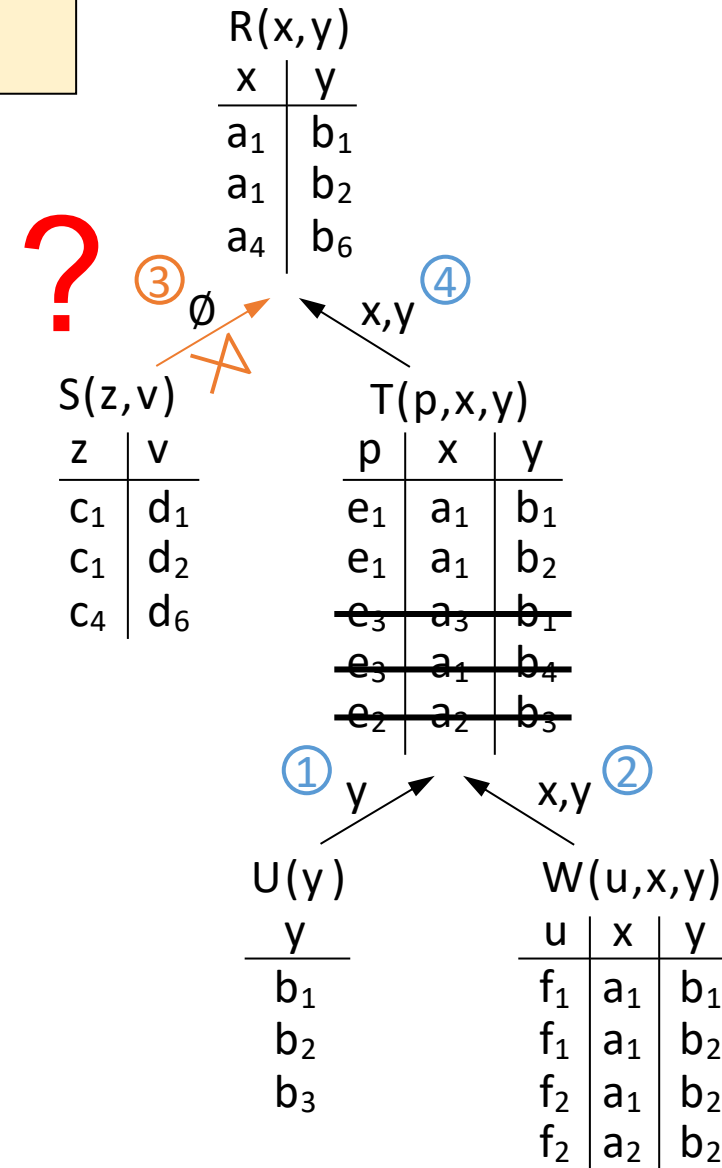


Yannakakis Algorithm example: 1st pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order

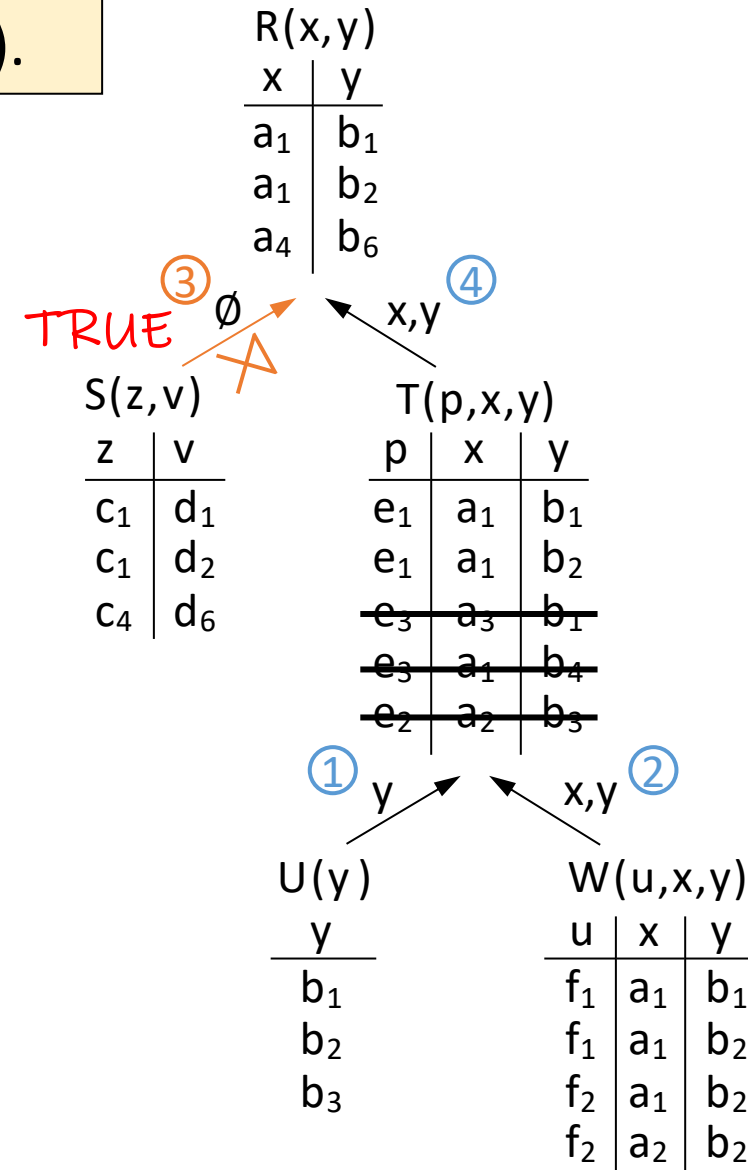


Yannakakis Algorithm example: 1st pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order

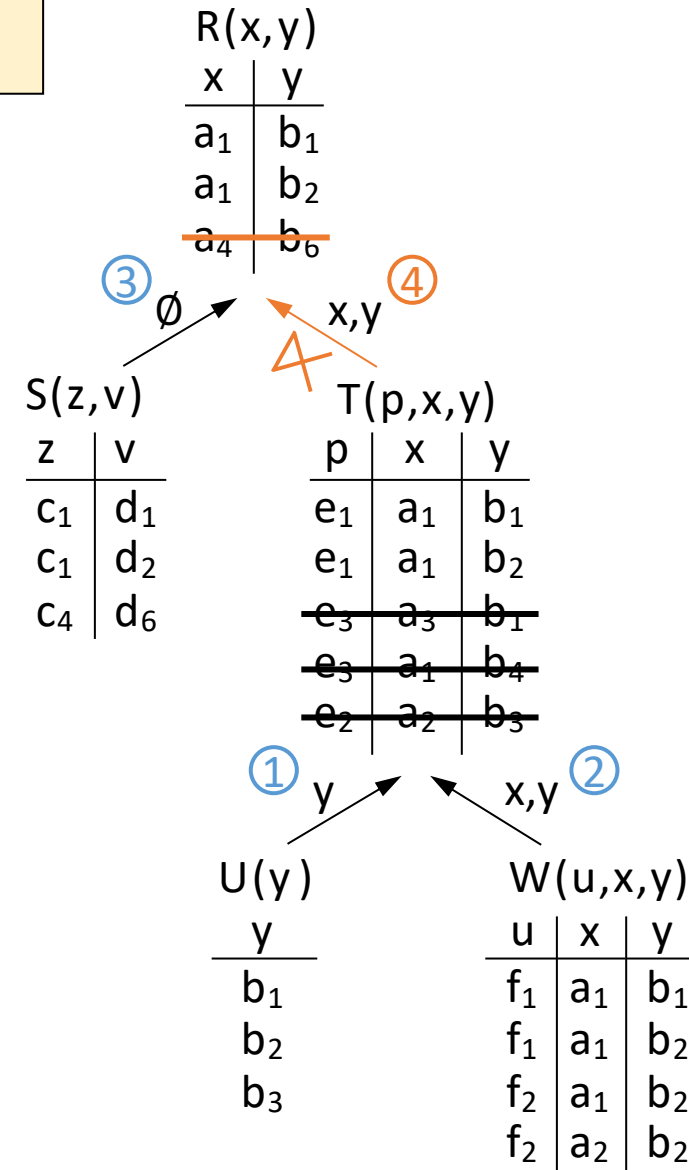


Yannakakis Algorithm example: 1st pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \times (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order



Yannakakis Algorithm example: 1st pass over the data

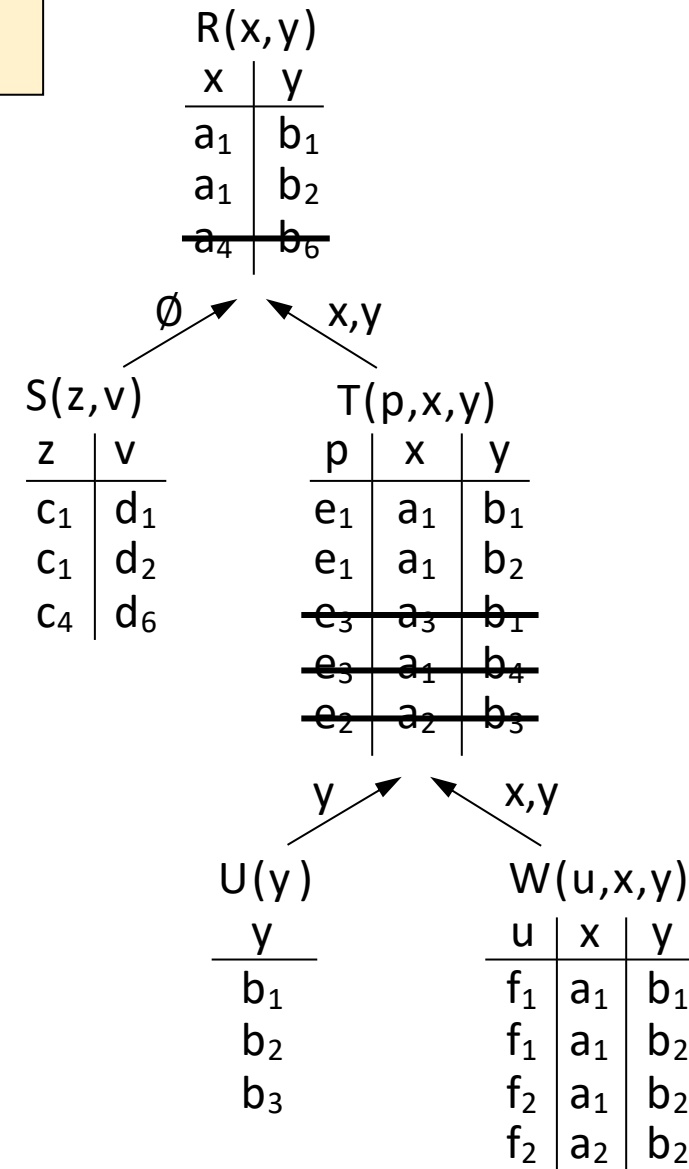
$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order

Notice that at the end of the 1st pass, the table R at the root does not contain any more dangling tuples; it is completely reduced.

In other words, with a sequence of only local updates, we have accumulated at the root all necessary information to answer the Boolean query.

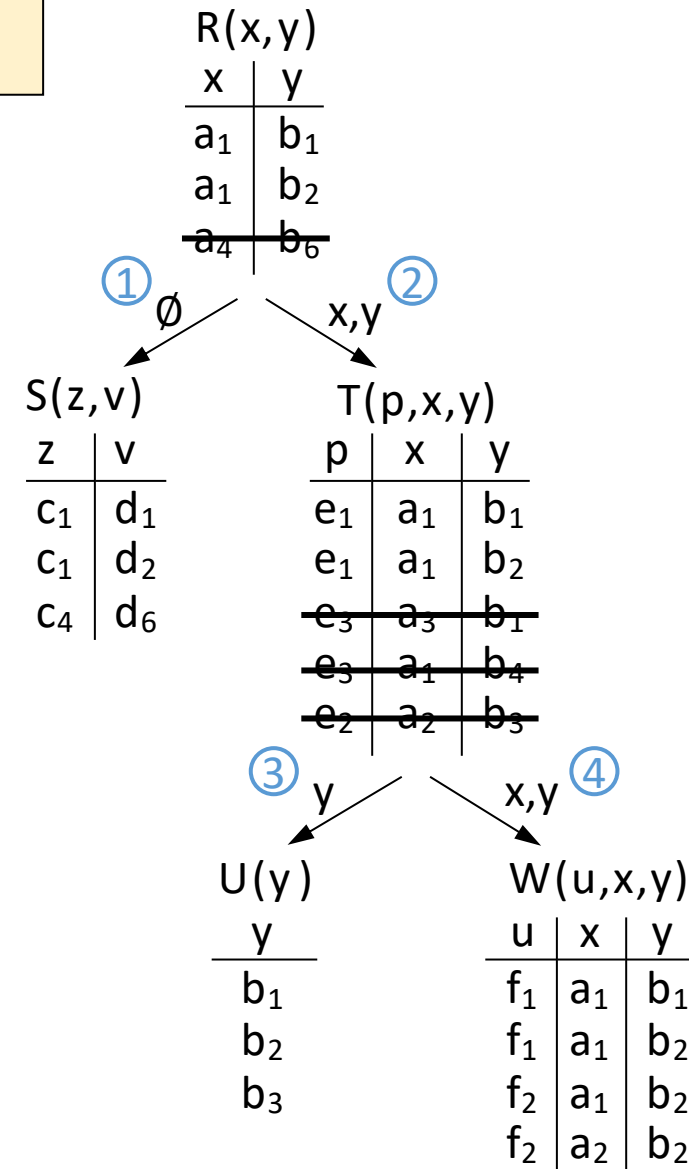


Yannakakis Algorithm example: 2nd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**

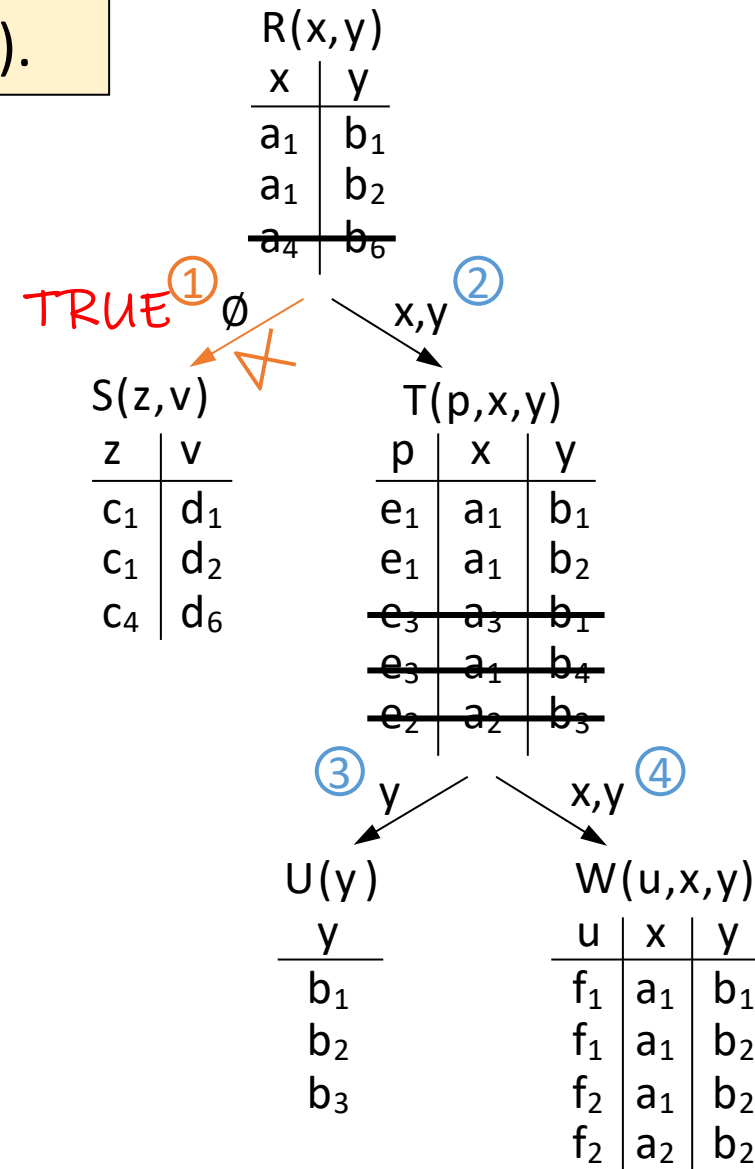


Yannakakis Algorithm example: 2nd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**

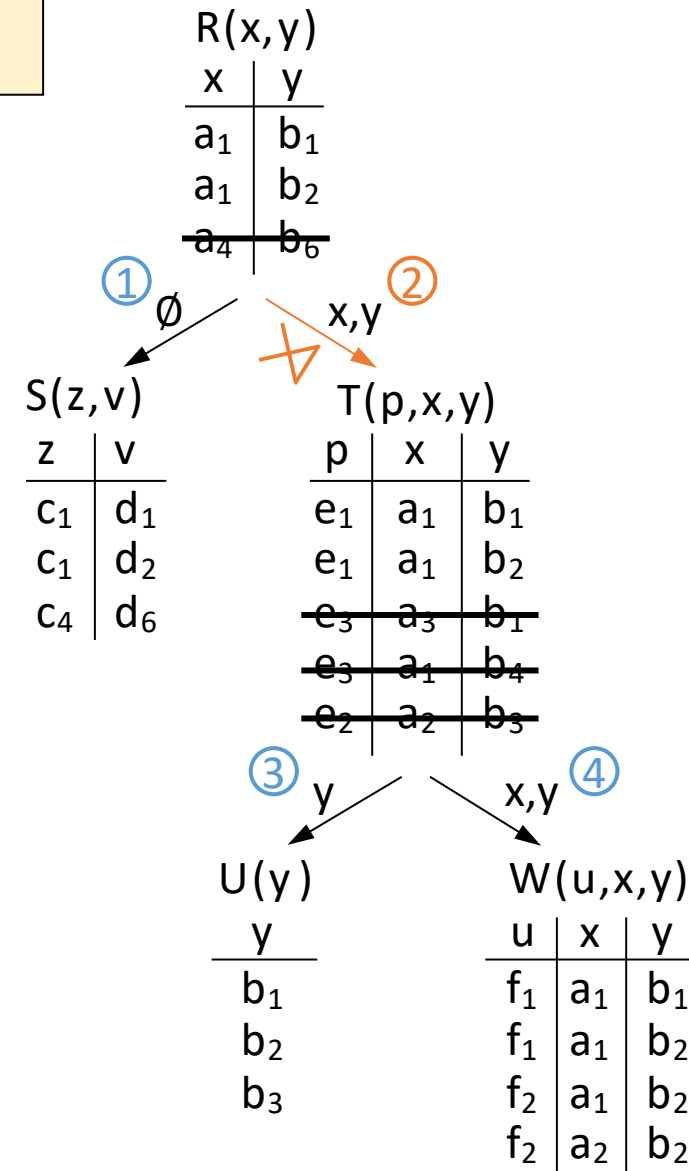


Yannakakis Algorithm example: 2nd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**

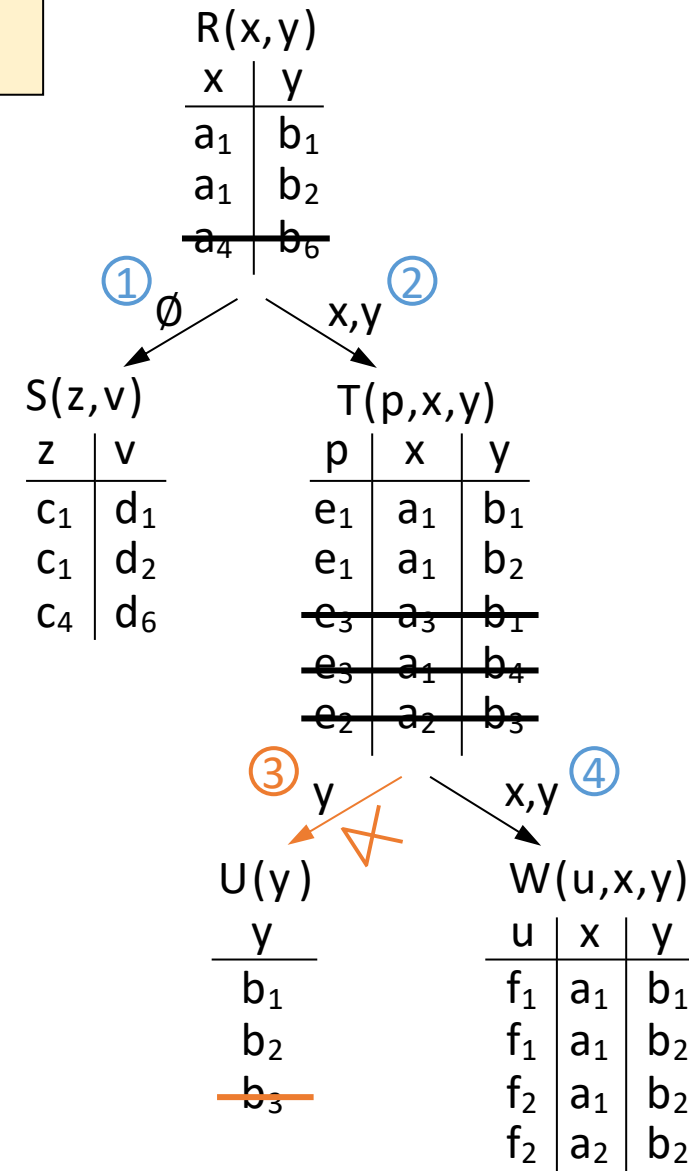


Yannakakis Algorithm example: 2nd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**

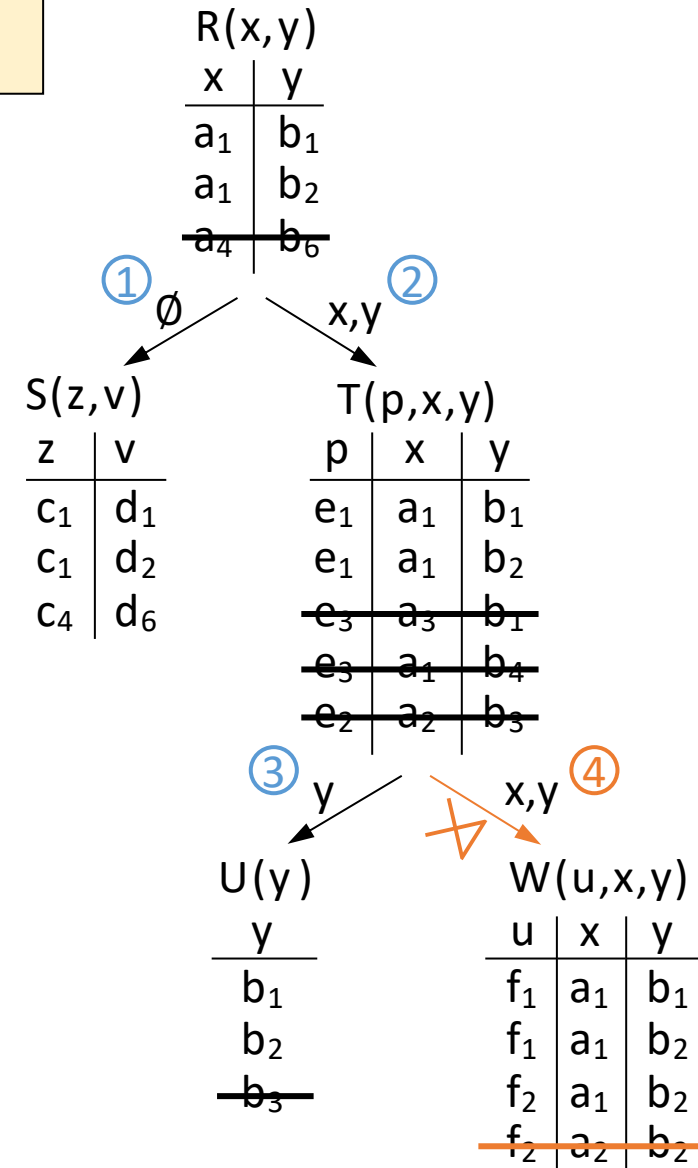


Yannakakis Algorithm example: 2nd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**



Yannakakis Algorithm example: 2nd pass over the data

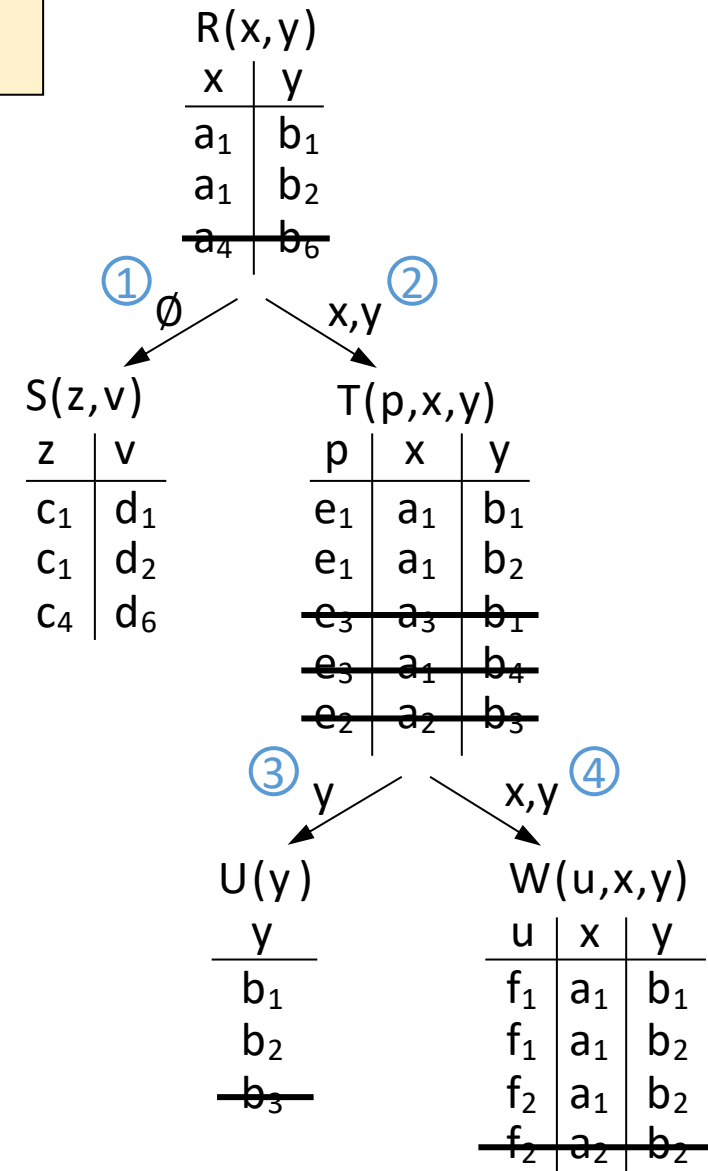
$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- **Top-down semi-join propagation from root to leaves in some topological order**

Notice that at the end of the second pass, all tables are reduced; no table contains any more dangling tuples.

In other words, *every* table now "knows" whether the Boolean version of the query is true.



Yannakakis Algorithm example: 3rd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

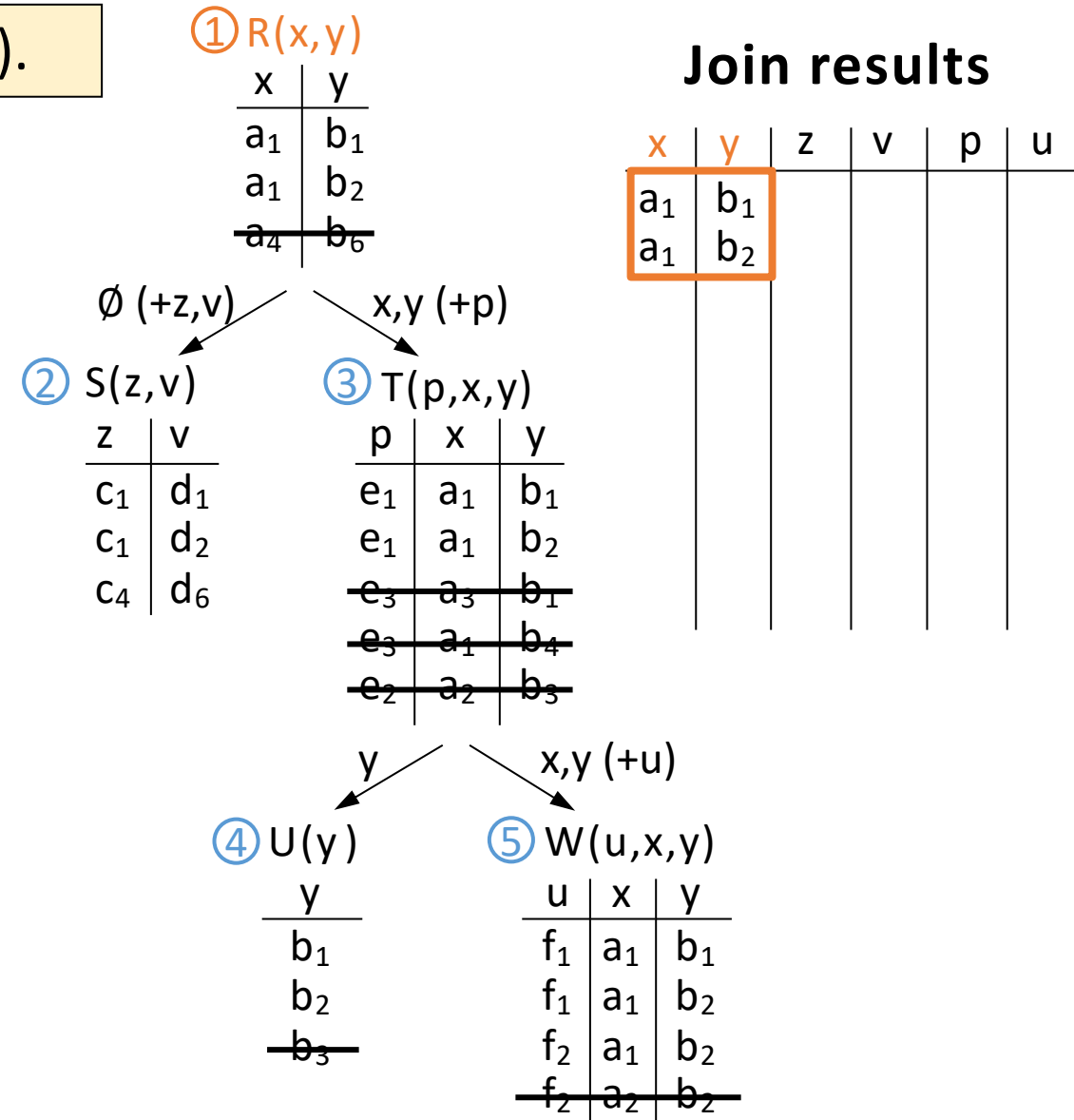
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. **Join phase** \bowtie (compute results) in $O(r)$ ← *Output*

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

Notice how with every join, the join result can never decrease in size!



Yannakakis Algorithm example: 3rd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

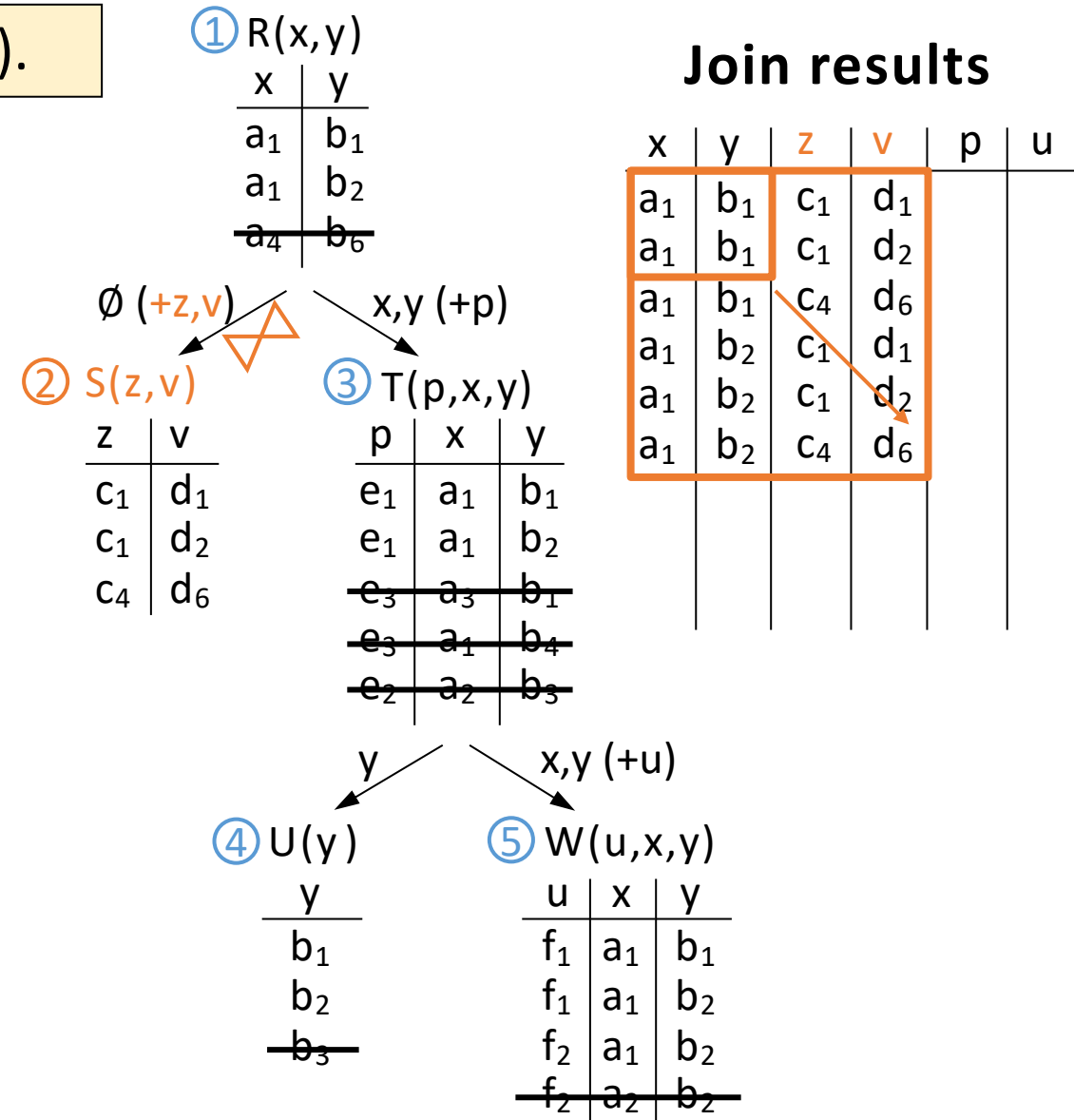
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. **Join phase** \bowtie (compute results) in $O(r)$ ← *Output*

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

Notice how with every join, the join result can never decrease in size!



Yannakakis Algorithm example: 3rd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

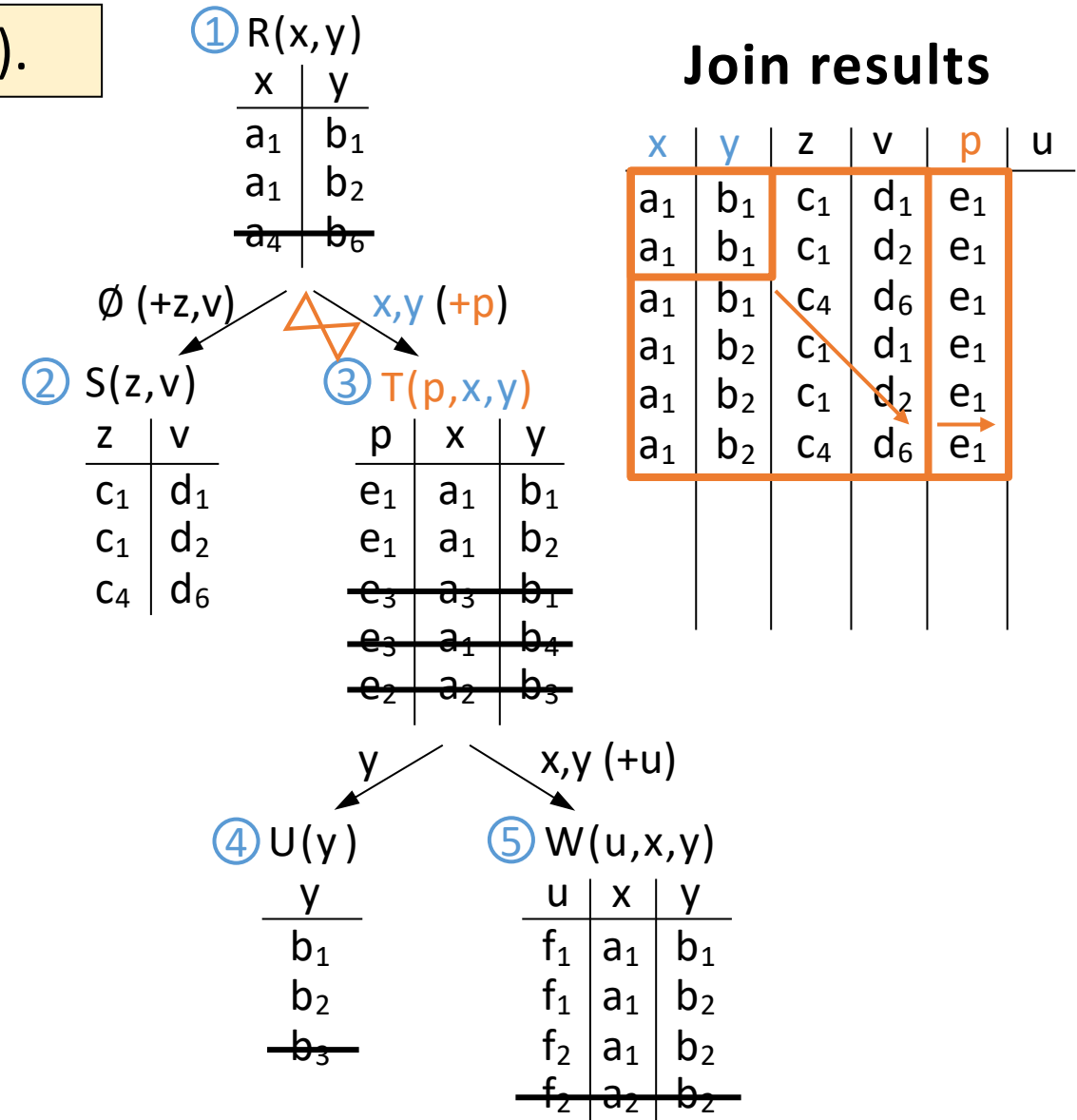
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Join phase \bowtie (compute results) in $O(r)$

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

Notice how with every join, the join result can never decrease in size!



Yannakakis Algorithm example: 3rd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

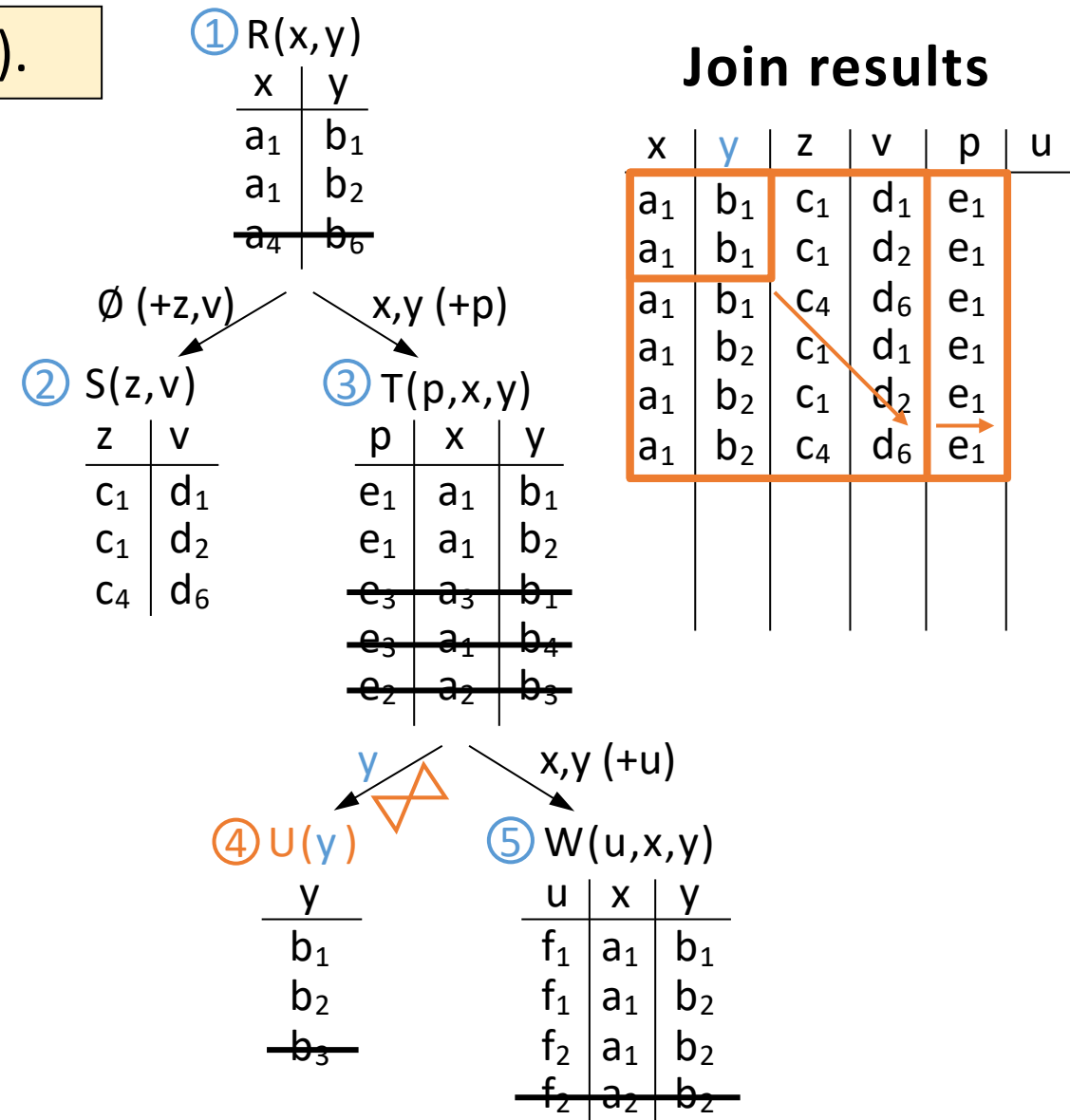
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Join phase \bowtie (compute results) in $O(r)$

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

Notice how with every join, the join result can never decrease in size!



Yannakakis Algorithm example: 3rd pass over the data

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

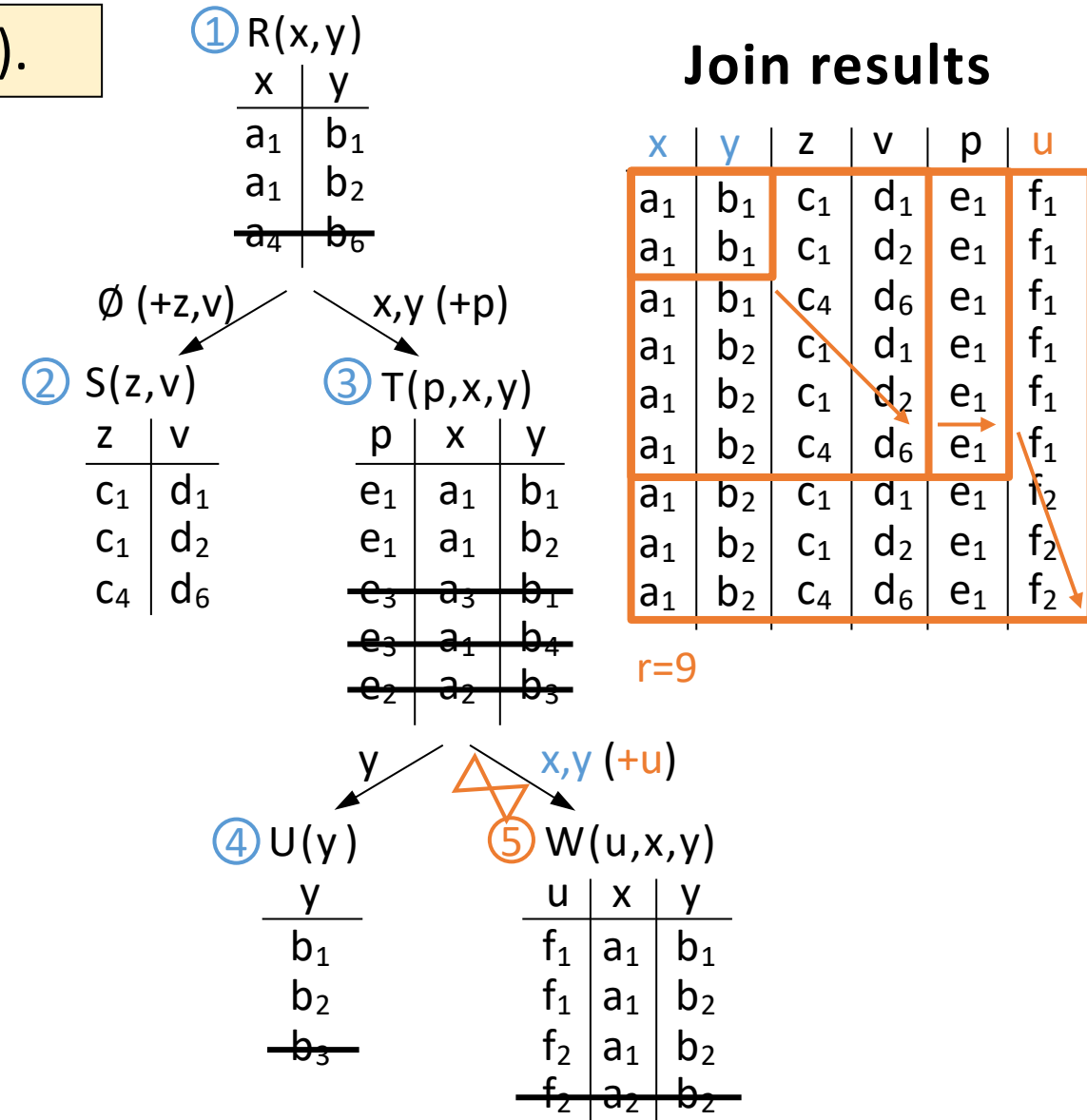
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Join phase \bowtie (compute results) in $O(r)$

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

Notice how with every join, the join result can never decrease in size!



Yannakakis Algorithm example: summary



$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

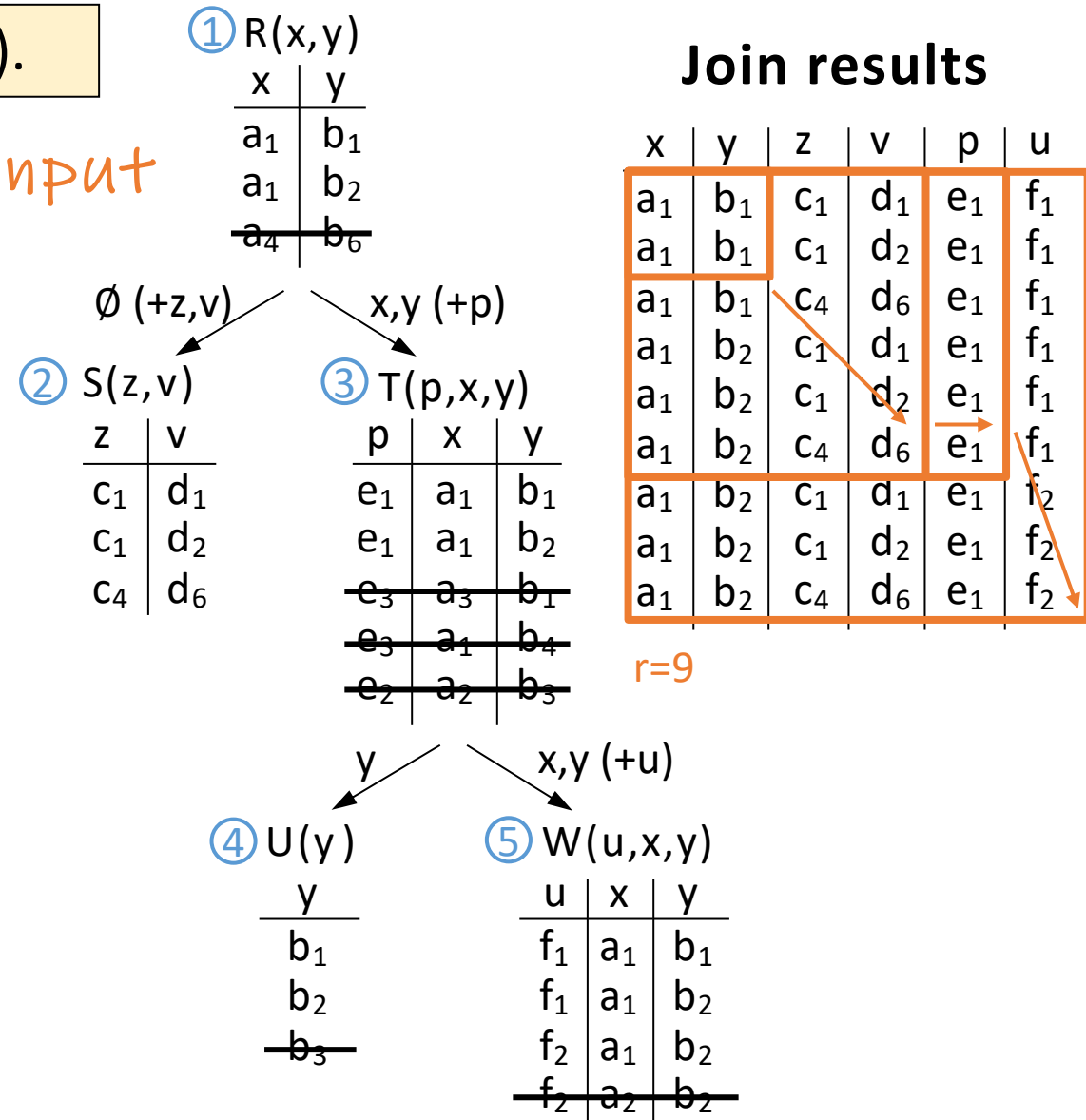
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$ \leftarrow *Input*

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. *Join phase* \bowtie (compute results) in $O(r)$ \leftarrow *Output*

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊

*Deciding $Q(D) \neq \emptyset$ is in $O(n)$
Computing $Q(D)$ is in $O(n+r)$*



Deciding ACQs Efficiently [Yannakakis'81]

- Non-Serial Dynamic Programming (NSDP) algorithm over a **join tree** $T = (V, E)$ of a query Q , given database instance D
- Decide Boolean variant $Q(D) = \emptyset$ as follows:
 - Pick a root and assign to each $R_j \in V$ the corresponding relation R_j^D of D
 - In a bottom-up reverse topological order of T : compute semijoins of R_j^D
 - If the thus reduced relation at root is empty, then $Q(D) = \emptyset$, else $Q(D) \neq \emptyset$.
- Theorem:
 - For ACQs Q : Deciding $Q(D) = \emptyset$ is feasible in INPUT linear time.
 - Computing $Q(D)$ can be done in OUTPUT polynomial time.
 - For full queries (no projections) in OUTPUT linear time.

Recap topics 2 & 3

- T2:
 - The notions and complexity of query equivalence and containment
 - The Homomorphism Theorem
 - Minimization of conjunctive queries
- T3 so far: *Acyclic conjunctive queries*
 - Semi-join reductions, notions of hypergraph acyclicity
 - The Yannakakis algorithm
- T3 yet to be seen:
 - What do we do with cycles?
 - What about ranked retrieval?

Pointers to original work

- The original GYO algorithm was developed concurrently by Graham and Yu-Ozsoyoglu:
 - [Gra79] Graham. *On the universal relation*. Technical Report, University of Toronto, 1979.
 - [YO79] Yu, Ozsoyoglu. *An algorithm for tree-query membership of a distributed query*. COMPSAC, 1979.
<https://doi.org/10.1109/CMPSAC.1979.762509>
- Yannakakis. *Algorithms for acyclic database schemes*. VLDB 1981
<https://dl.acm.org/doi/10.5555/1286831.1286840>
- Bernstein, Chiu. *Using semi-joins to solve relational queries*. JACM 1981.
<https://doi.org/10.1145/322234.322238>
- Bernstein, Goodman. *Power of natural semi-joins*. SIAM J. 1981.
<https://doi.org/10.1137/0210059>
- Beeri, Fagin, Maier, Yannakakis. *On the desirability of acyclic database schemes*. JACM 1983.
<https://doi.org/10.1145/2402.322389>

Outline: T3-1: Acyclic conjunctive queries

- T3-1: Acyclic conjunctive queries
 - The semijoin operator
 - alpha-acyclic hypergraphs, join trees
 - GYO reduction
 - Full semi-join reductions
 - Yannakakis algorithm
 - Enumeration algorithms
- T3-2: Cyclic conjunctive queries

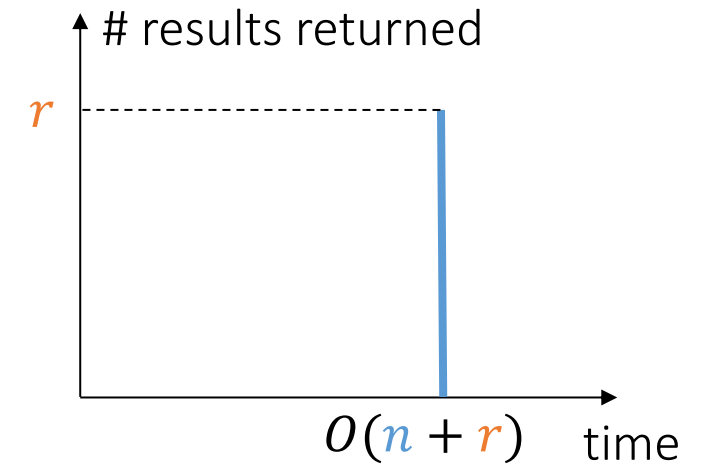
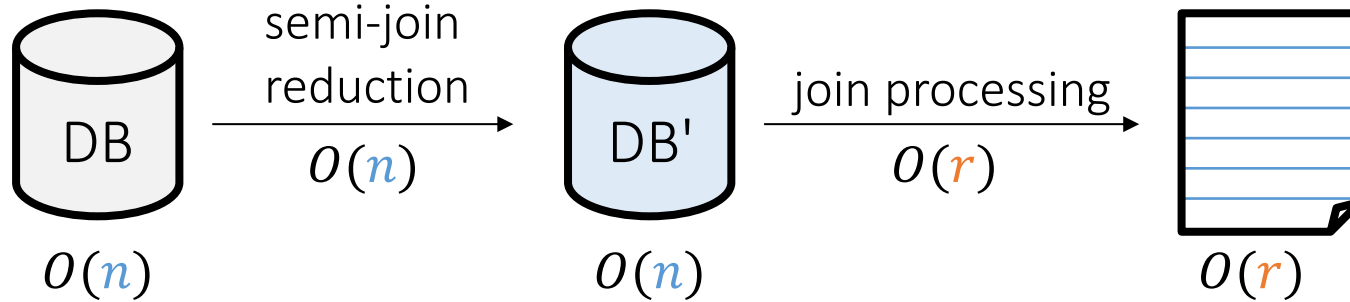
Several parts are an extended version of a tutorial from ICDE'22:

<https://www.youtube.com/watch?v=toi7ysuyRkw>

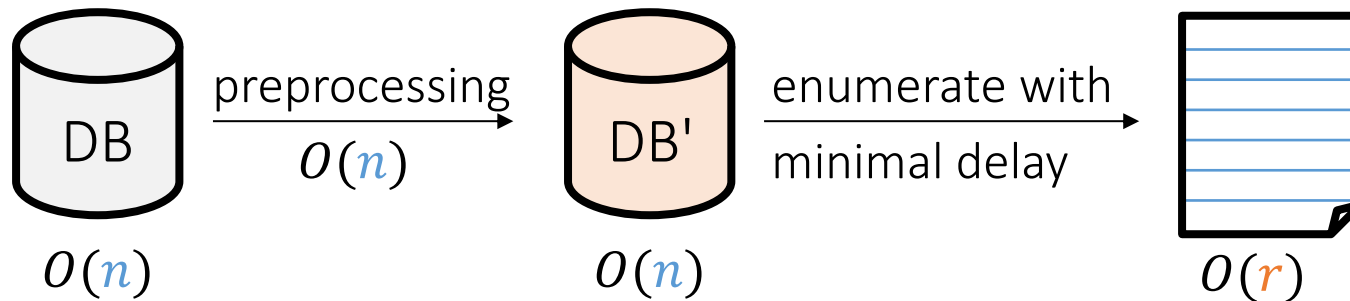
<https://northeastern-datalab.github.io/>

The enumeration framework

Standard Yannakakis framework for acyclic join processing

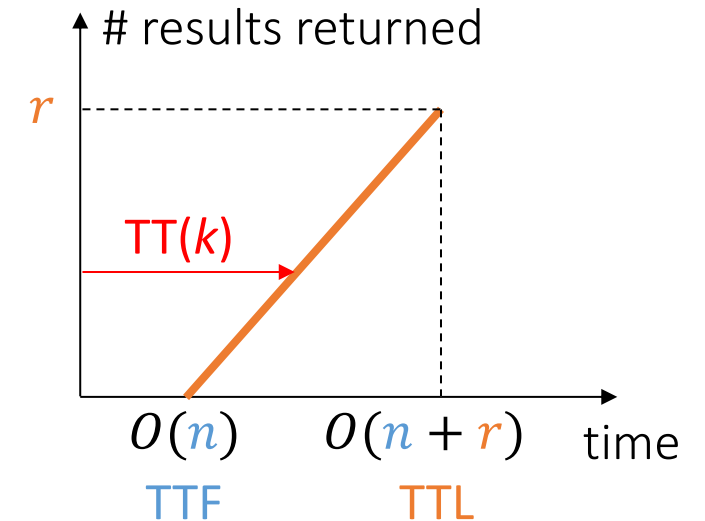


Enumeration framework for acyclic queries



TTF (Time-To-First)

TTL (Time-to-last)

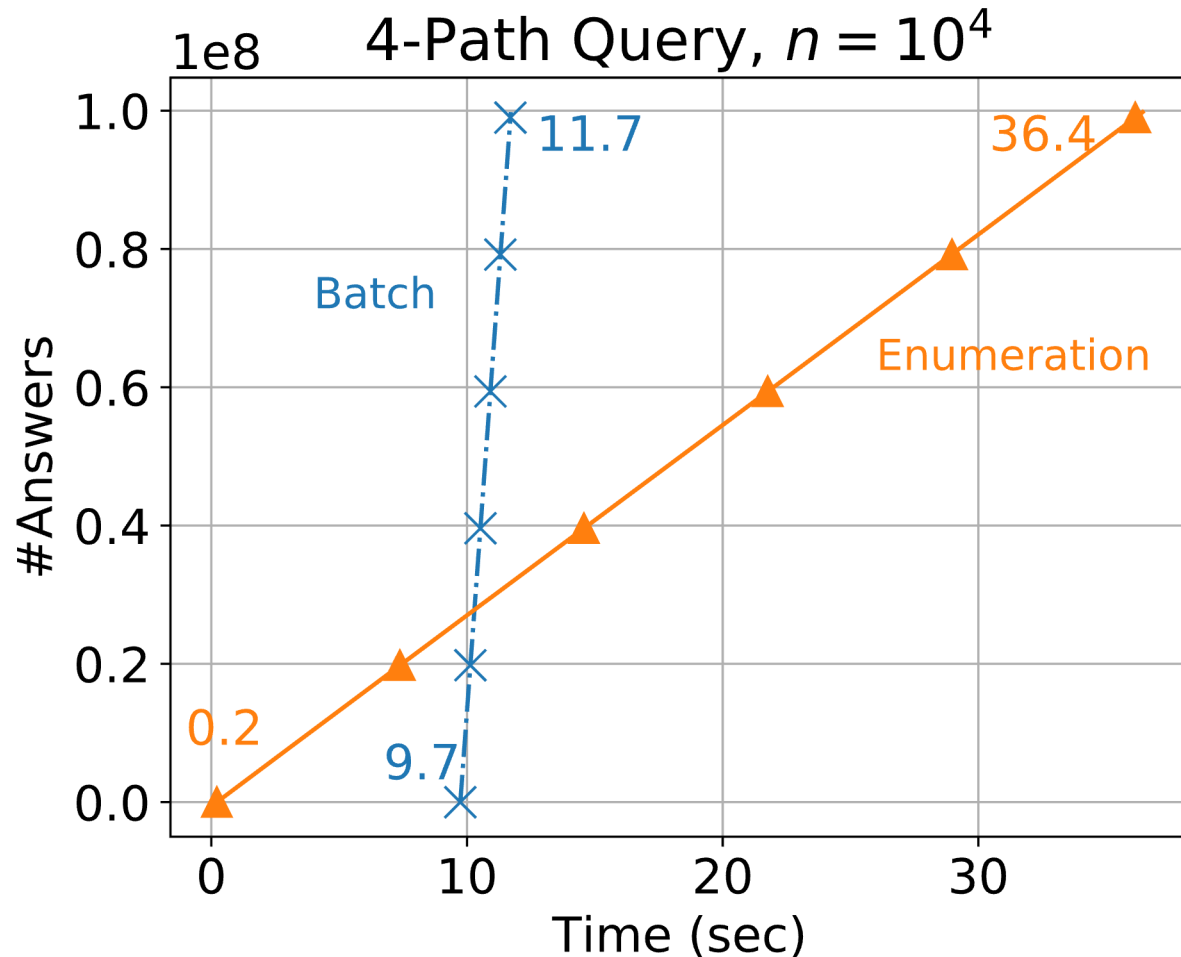


$r = O(n^{\rho^*})$ worst-case result size (AGM bound). ρ^* = fractional edge cover

Bagan, Durand, Grandjean. On acyclic conjunctive queries and constant delay enumeration. CSL 2007. https://doi.org/10.1007/978-3-540-74915-8_18

Towards Responsive DBMS. ICDE 2022 tutorial: <https://northeastern-datalab.github.io/responsive-dbms-tutorial>

Modified Yannakakis for output enumeration



- Standard Yannakakis:
 - Table-at-a-time / Breadth-first
 - After the semi-join reduction, Yannakakis visits each table once top-down, and at each stage increases the size of the answer set
- Enumeration:
 - Tuple-at-a-time / Depth-first
 - By a slight modification of Yannakakis and tuple by tuple

Yannakakis Algorithm: example from before

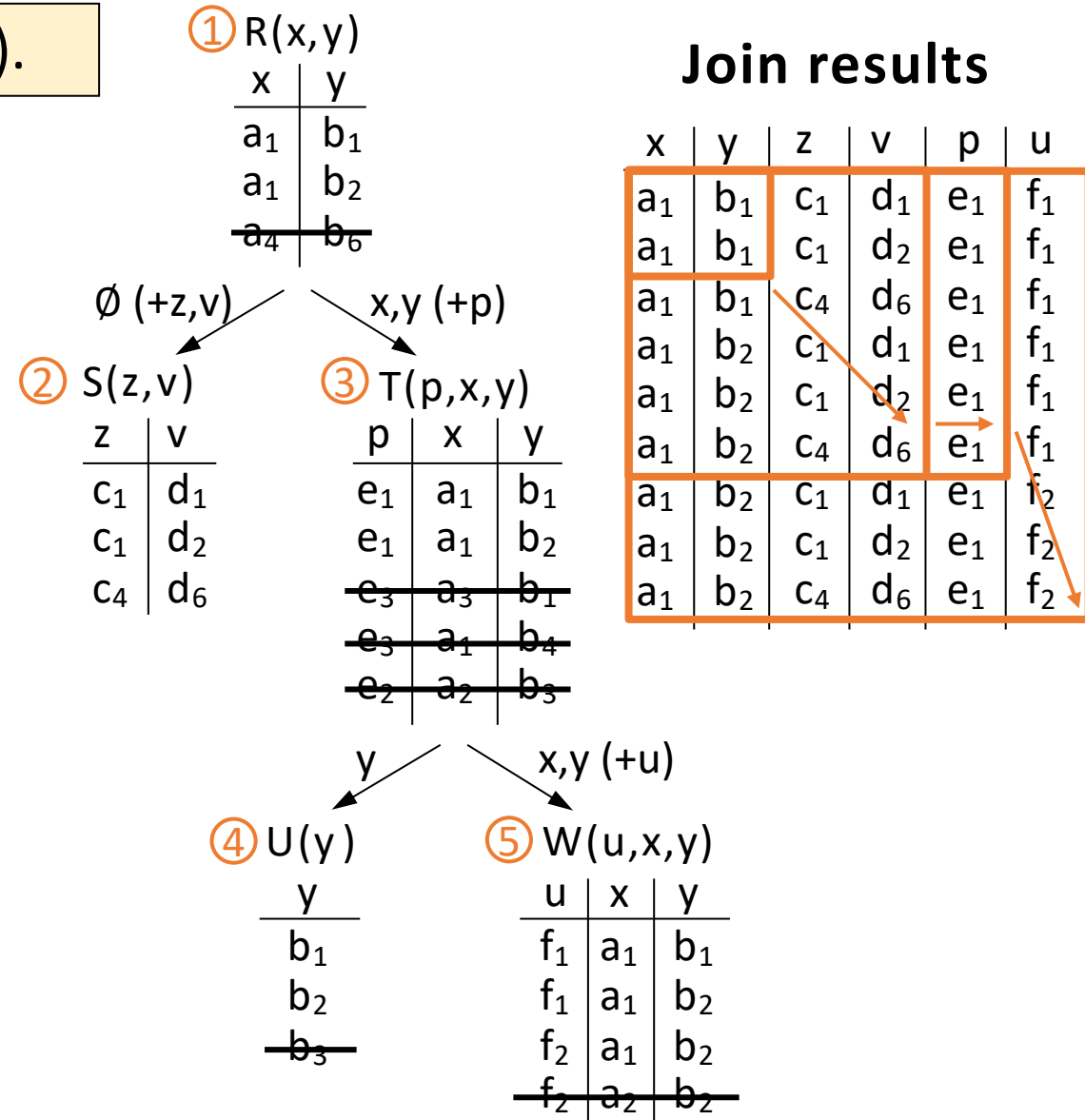
$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

1. Semi-join phase ✕ (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Join phase ✕ (compute results) in $O(r)$

- **Compute the results in a 2nd top-down (or 2nd bottom-up) traversal:**
 - This step can actually be combined with the earlier top-down traversal; thus two total passes (first from leaves, then from root) are actually enough 😊



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

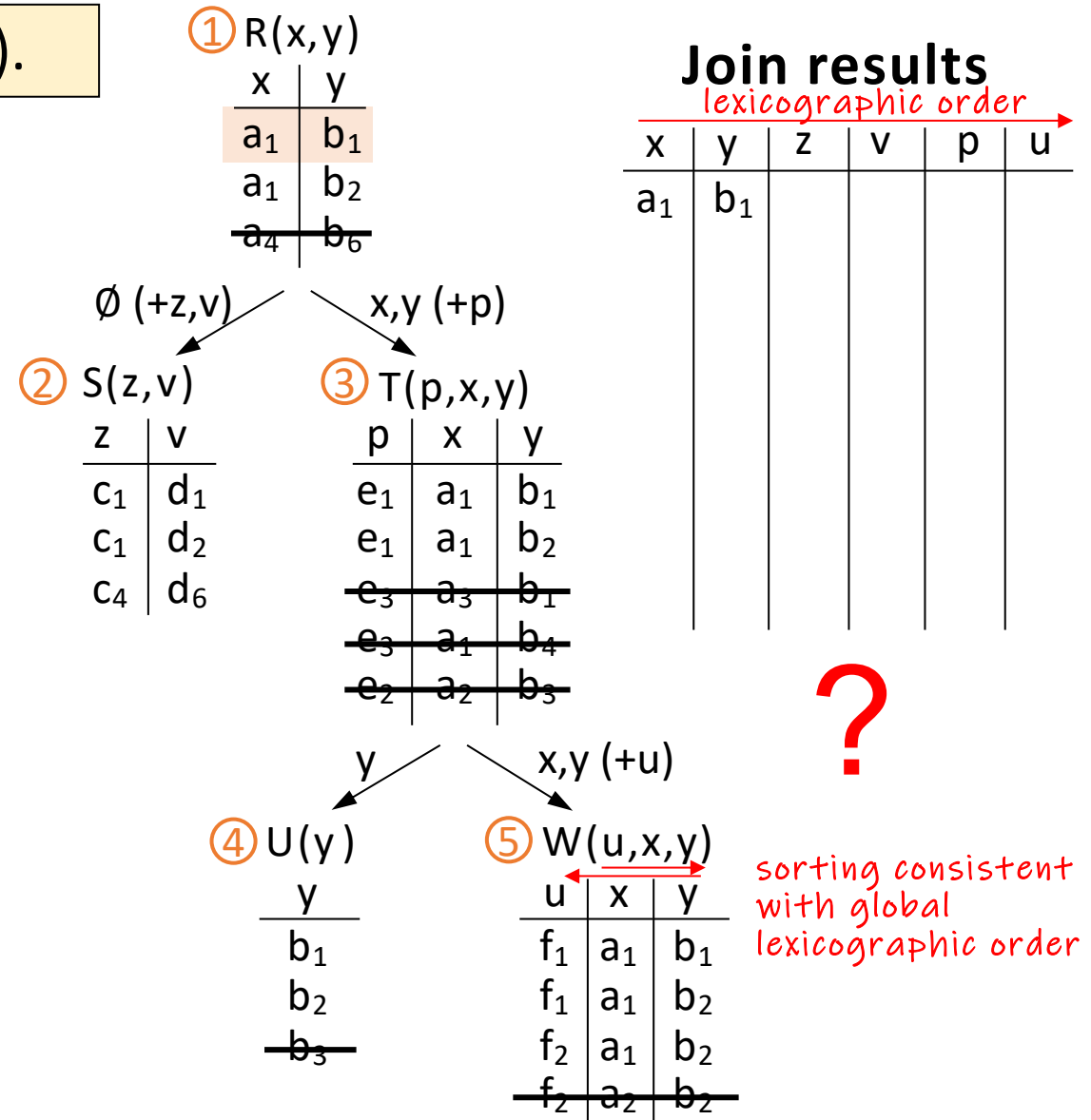
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

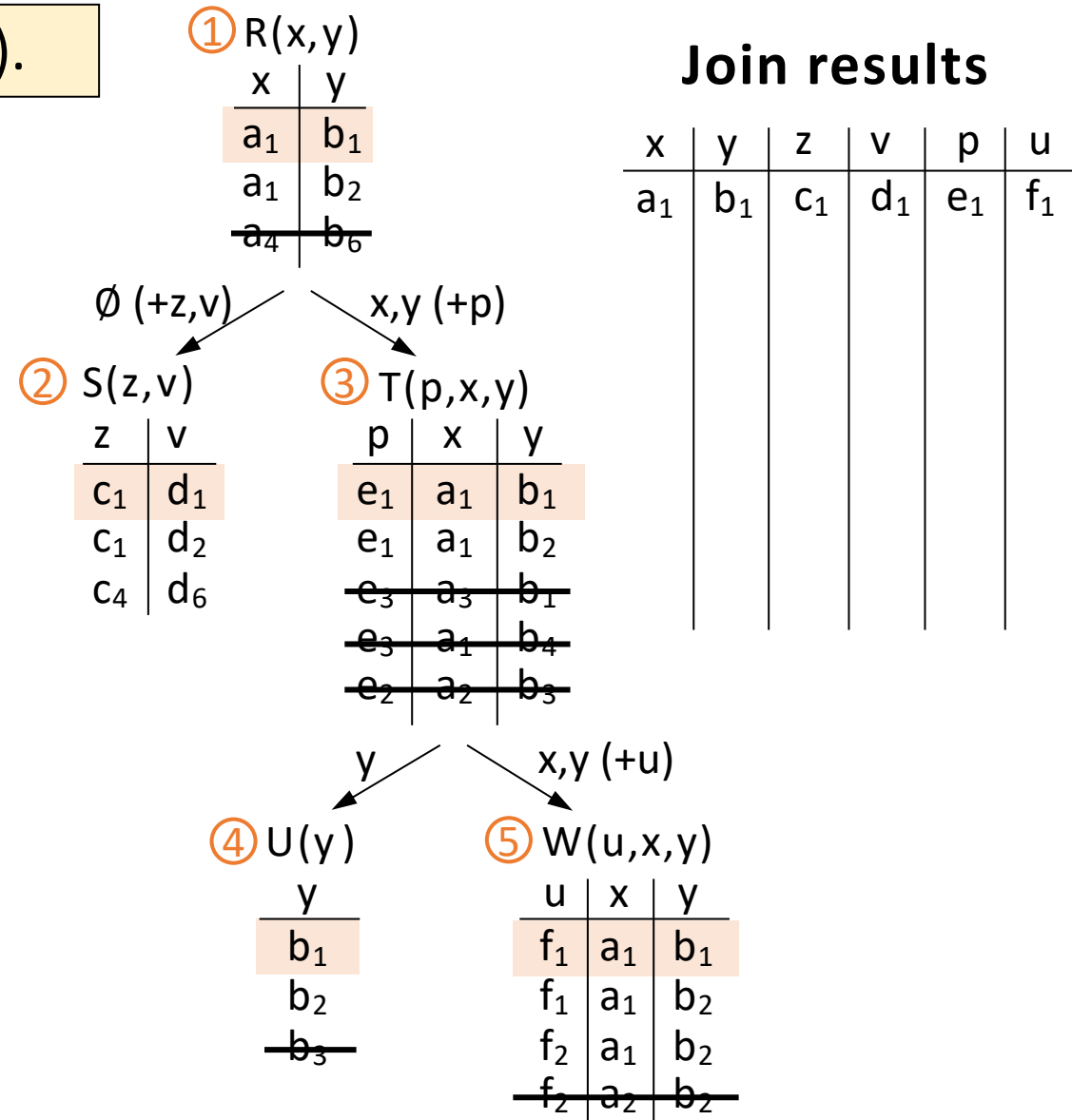
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

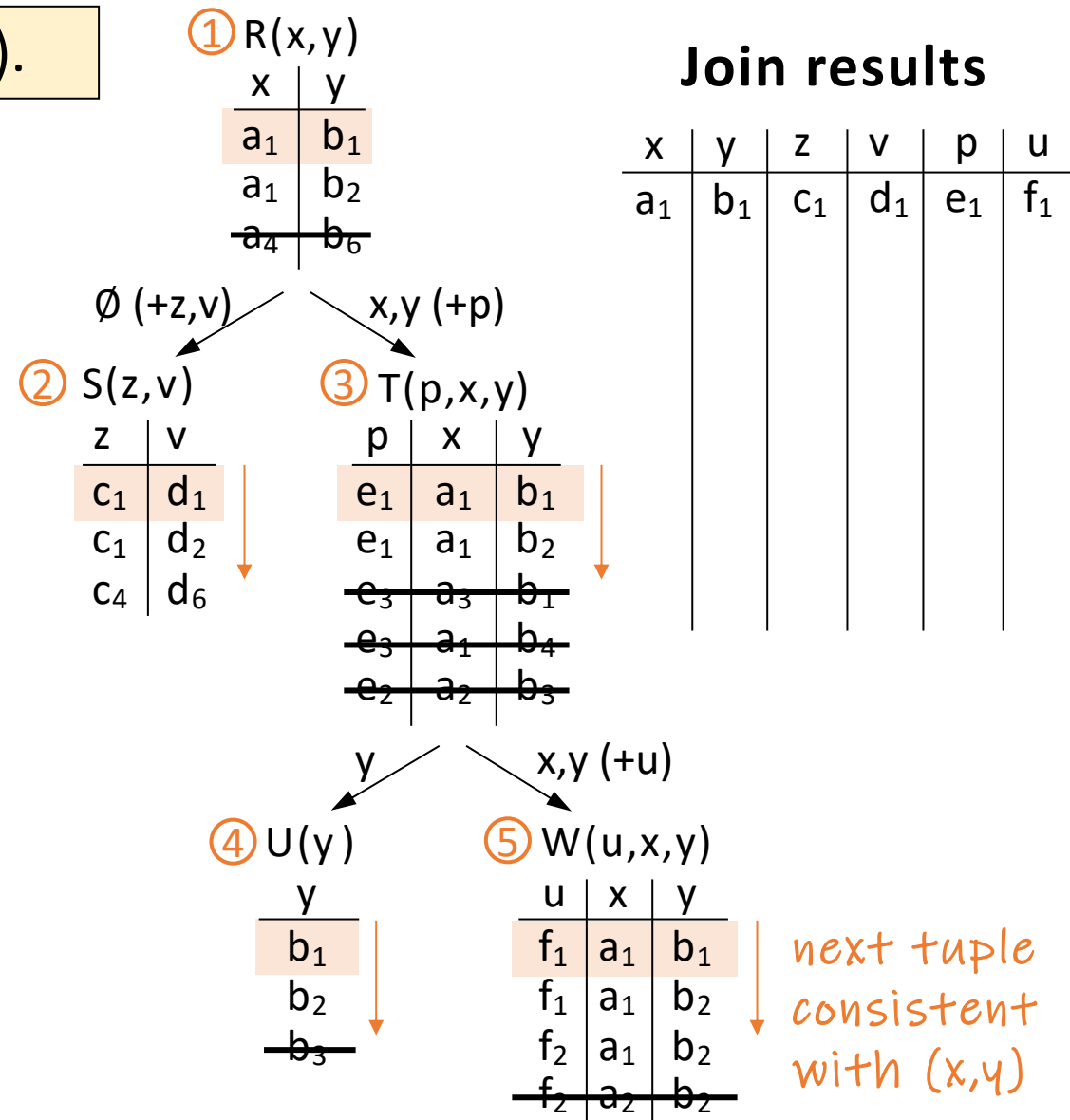
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\} + \{z,v\} + \{p\} + \emptyset + \{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

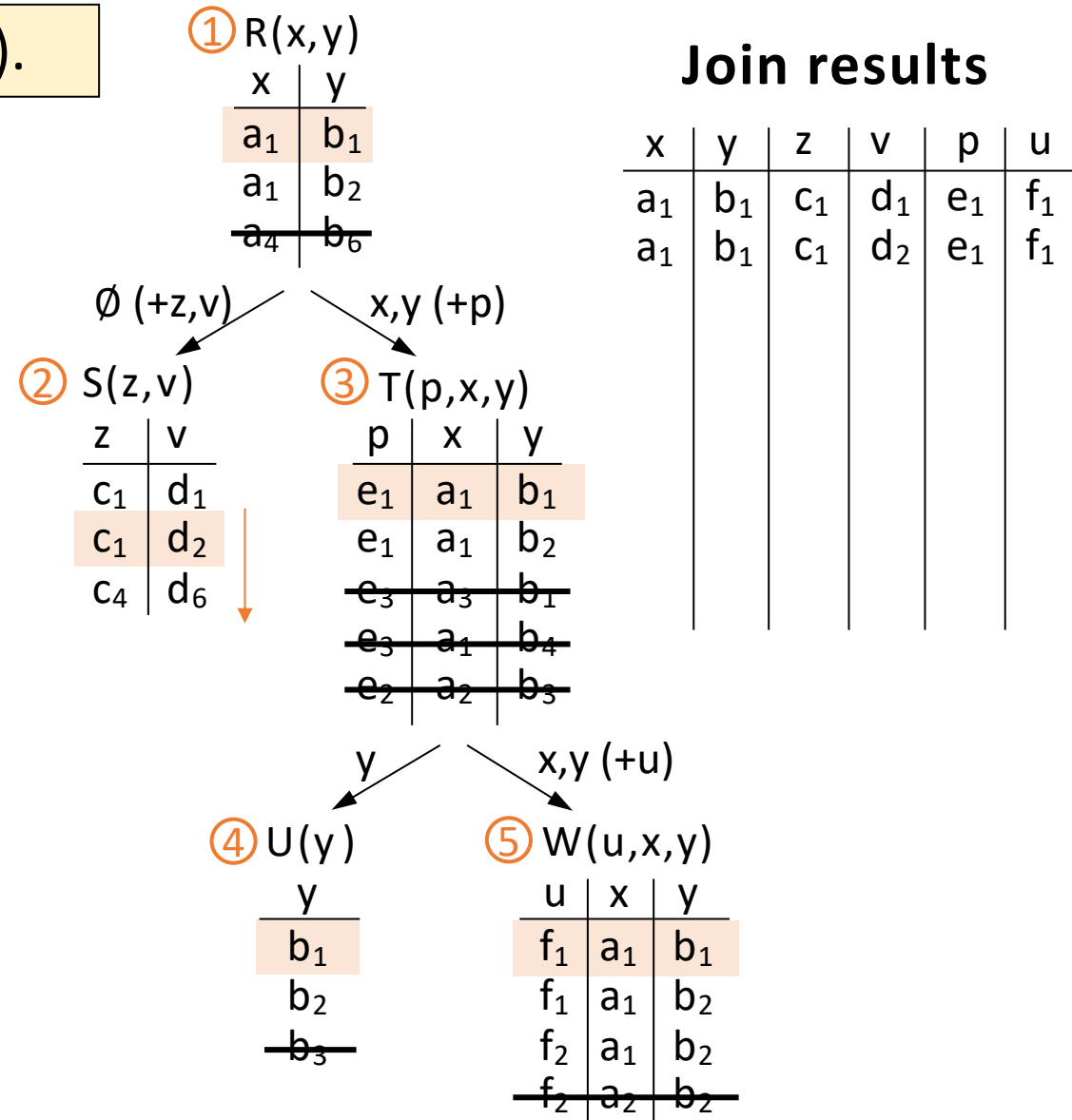
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

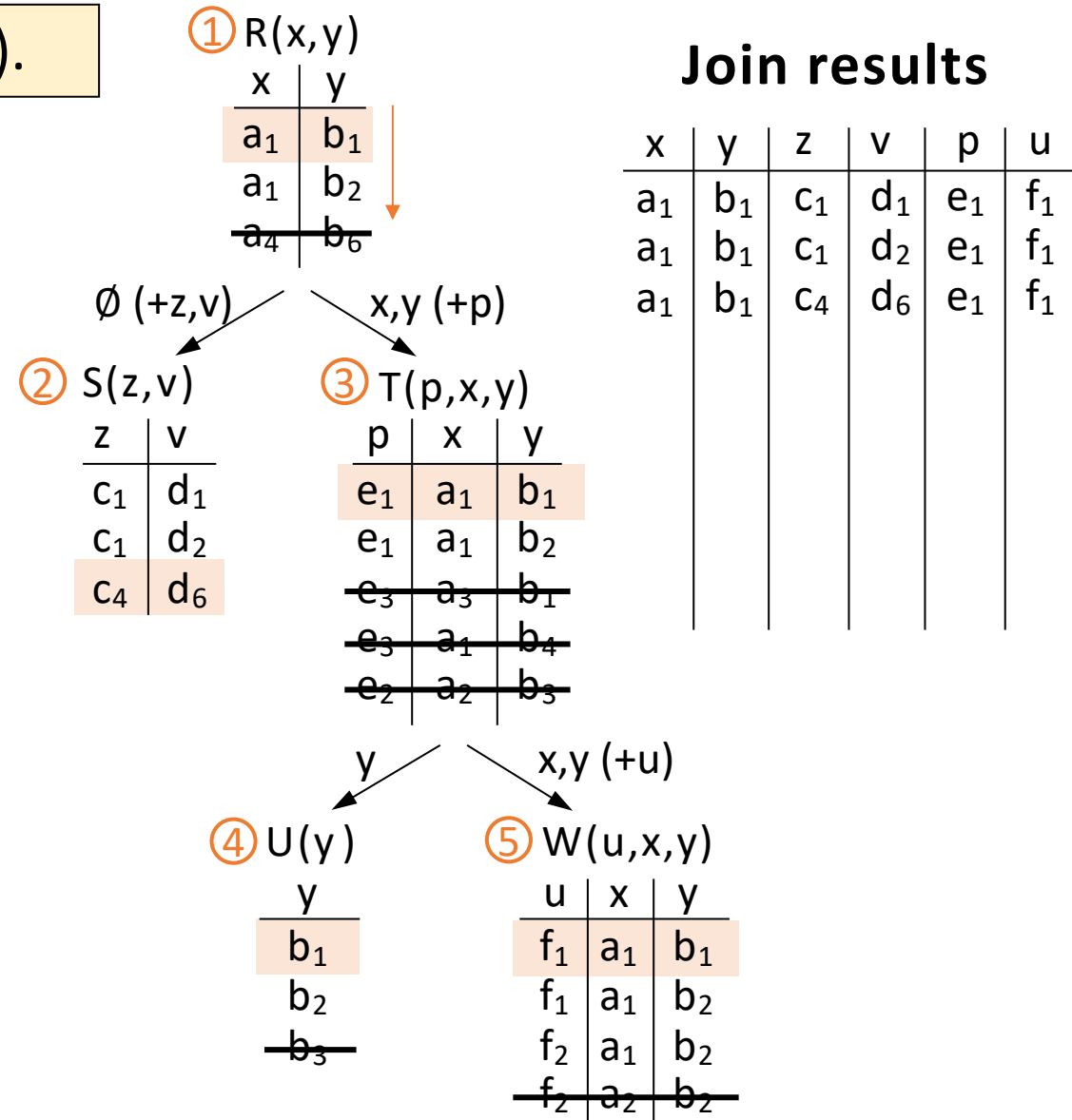
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

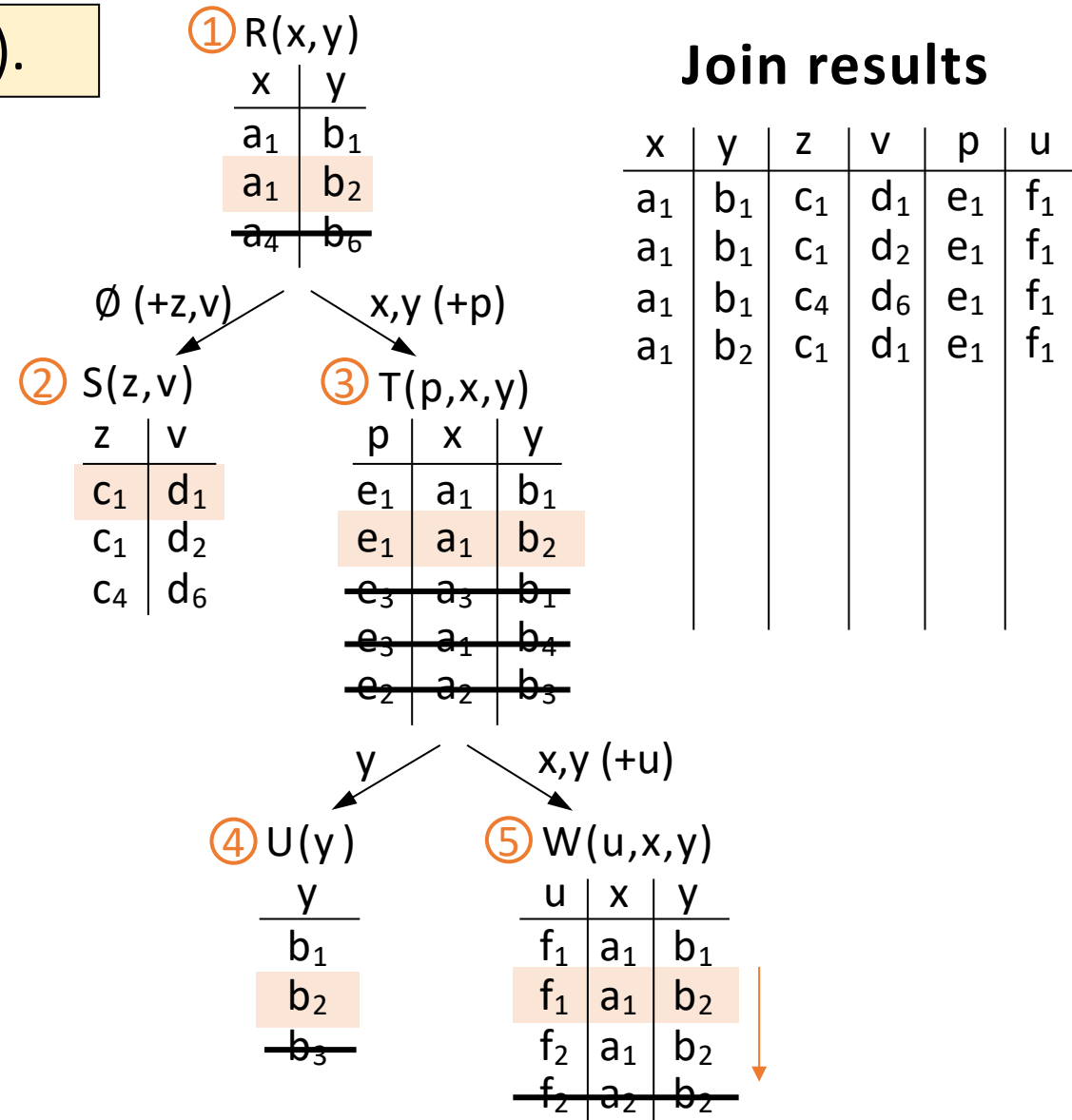
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

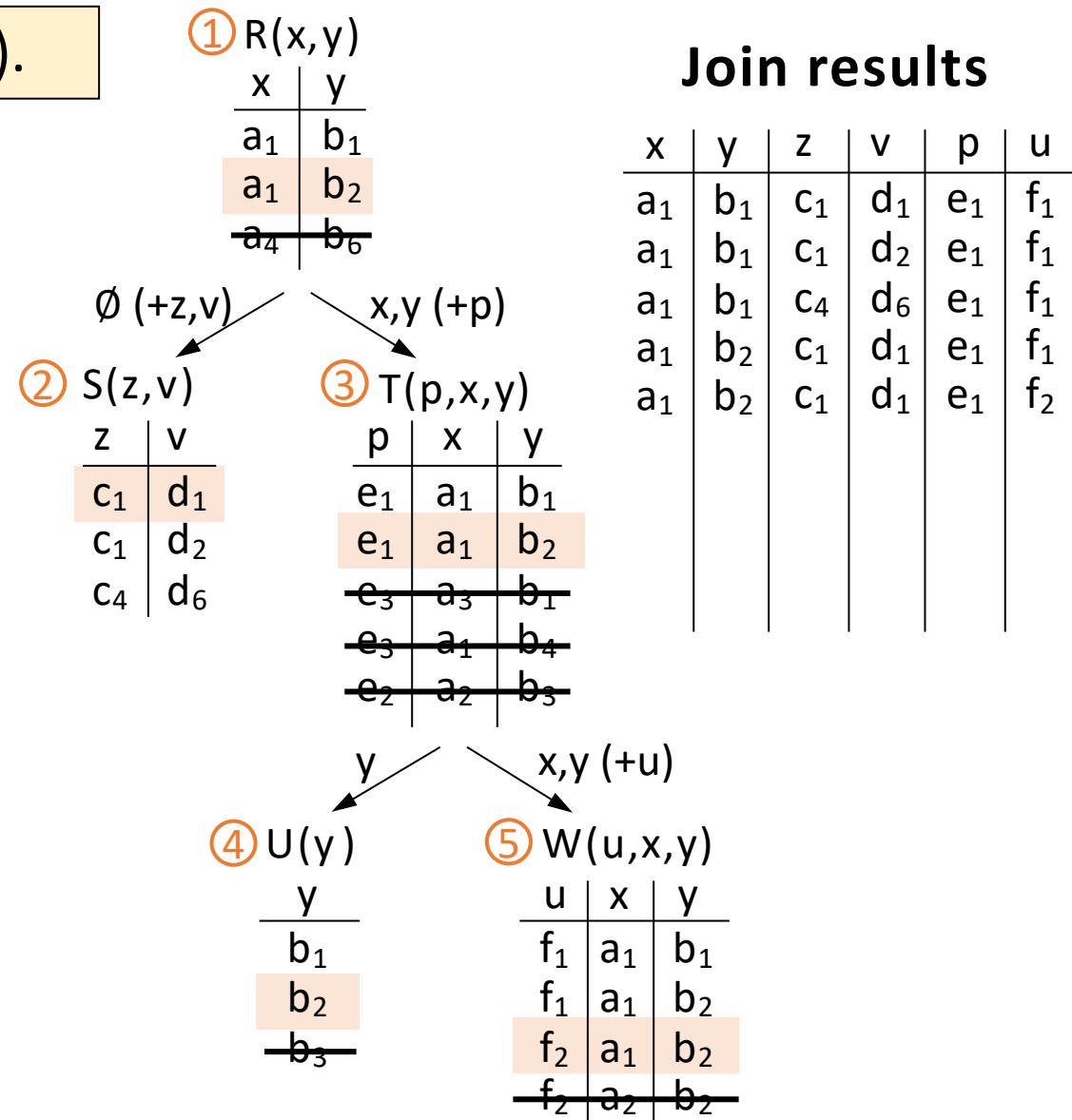
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

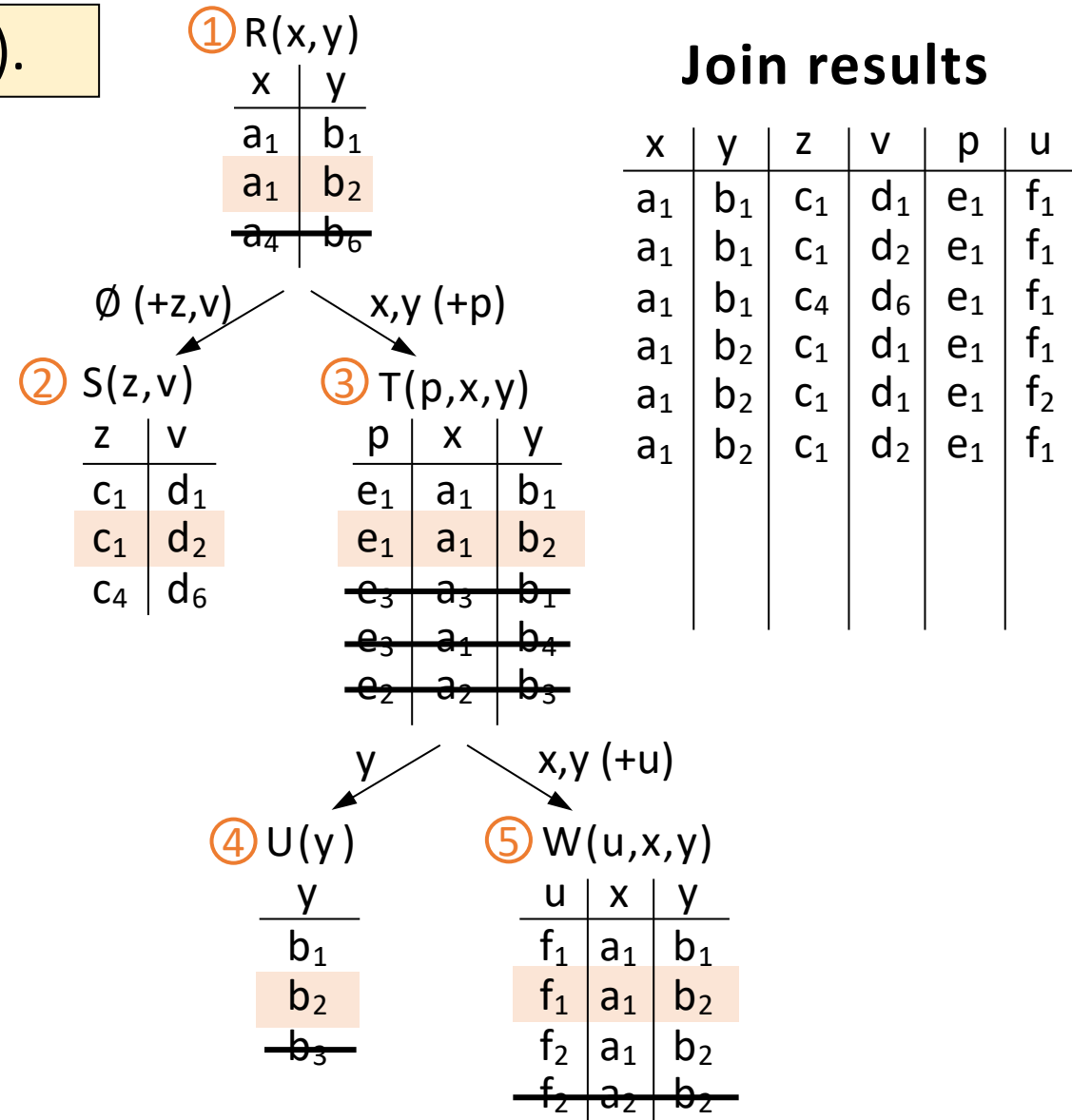
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

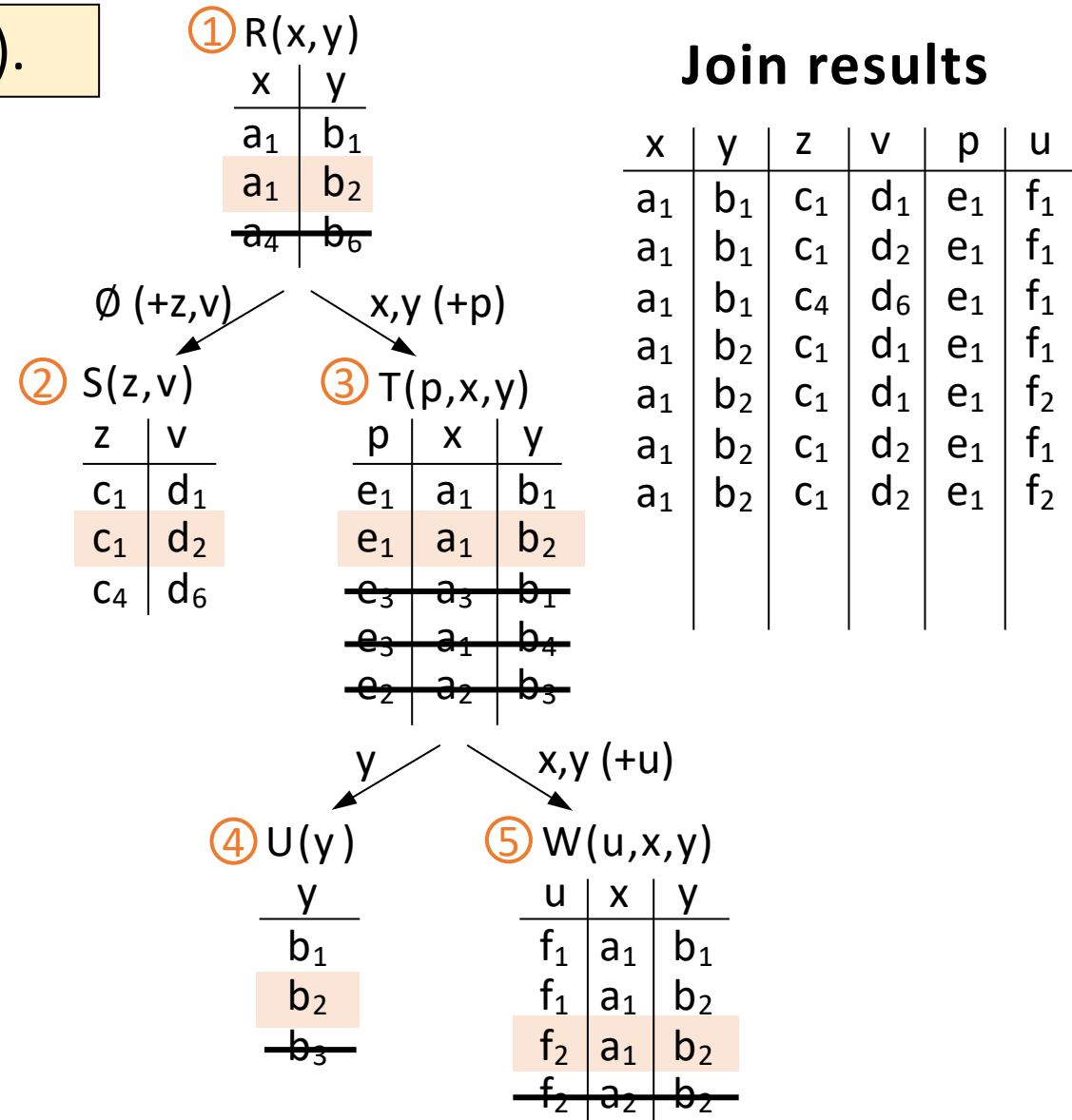
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

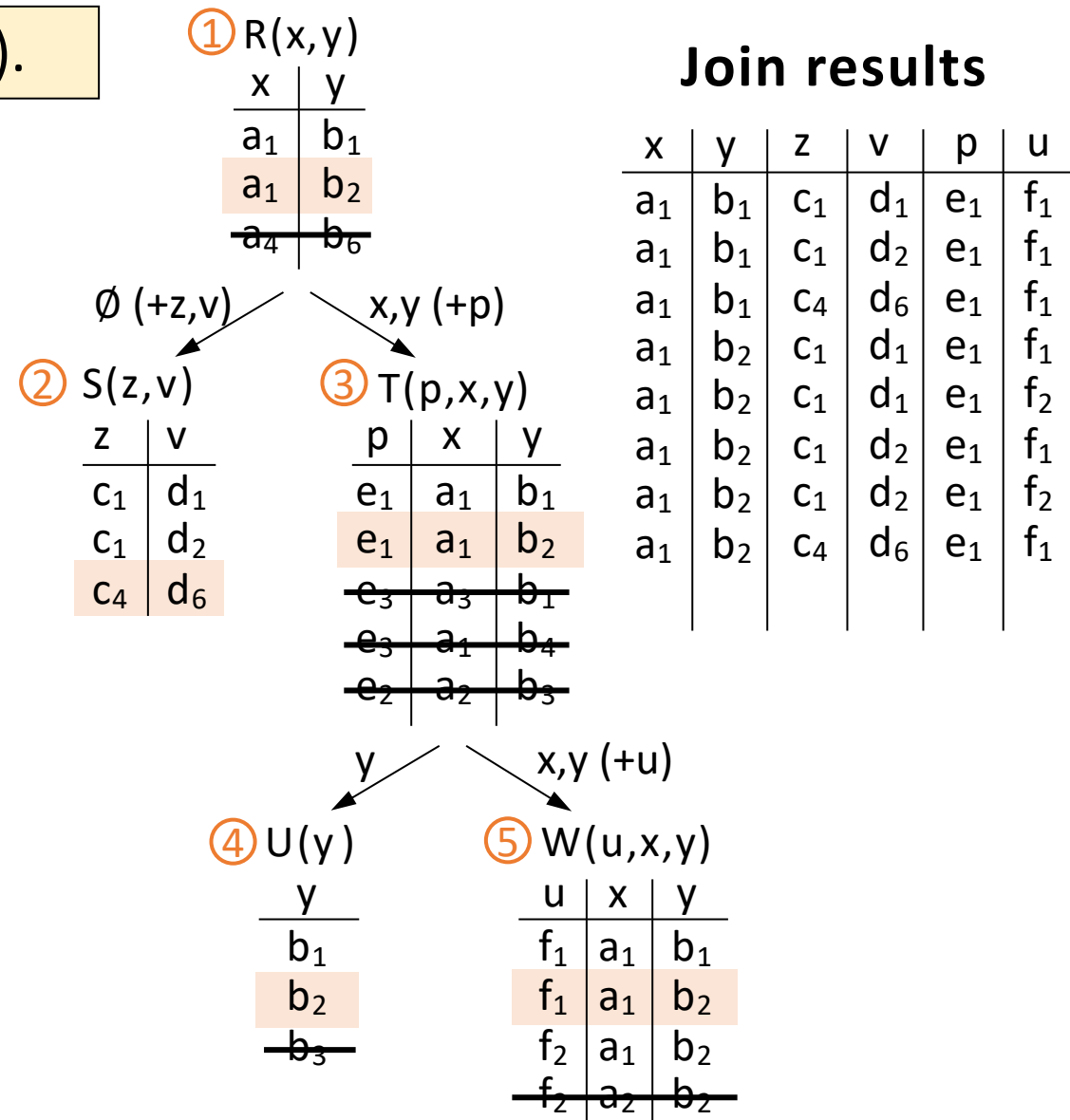
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis Algorithm: enumeration

$Q(x,y,z,v,p,u) :- R(x,y), S(z,v), T(p,x,y), U(y), W(u,x,y).$

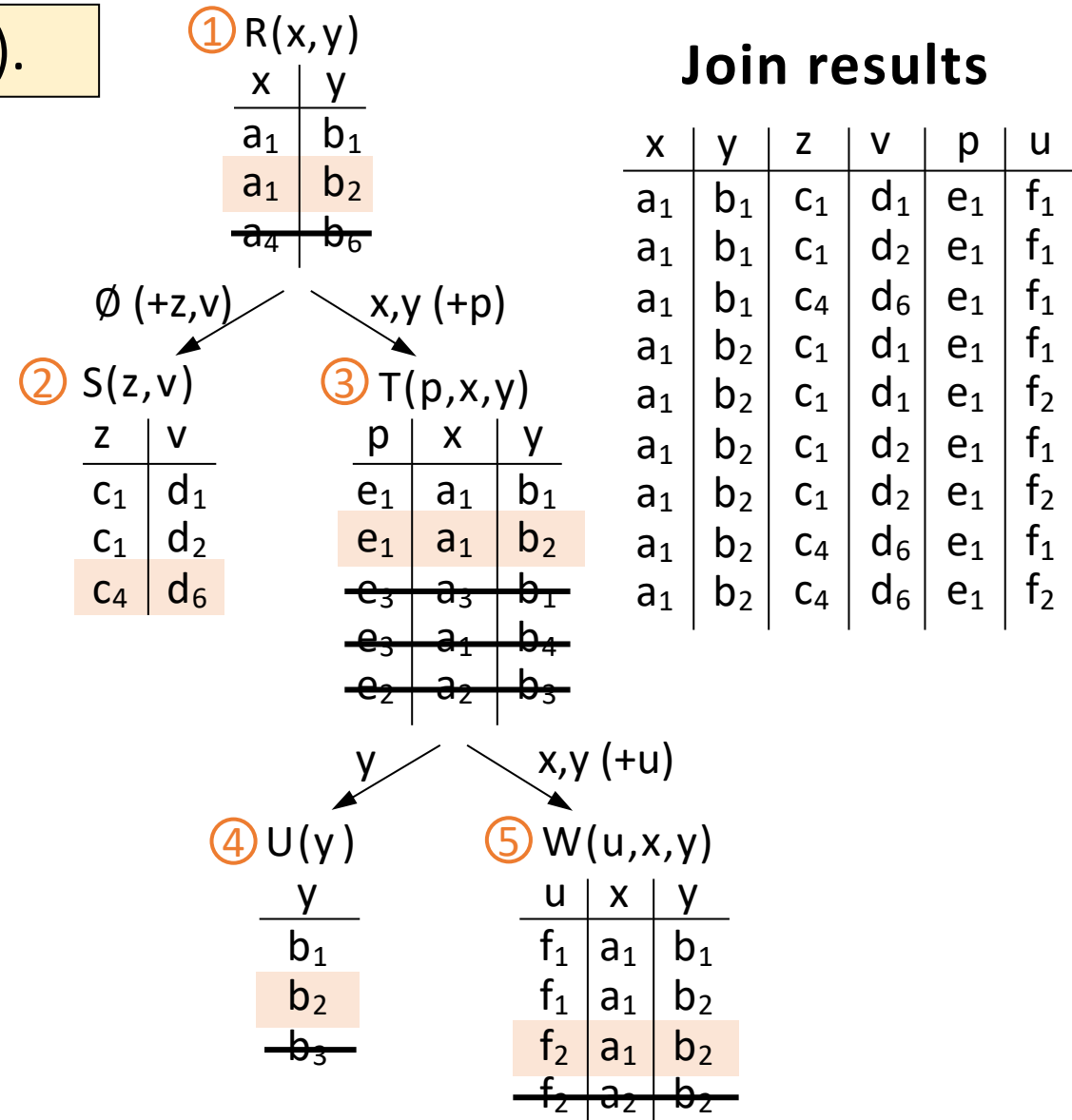
1. Semi-join phase \bowtie (remove dangling tuples) in $O(n)$

- Bottom-up semi-join propagation from leaves to root in some reverse topological order
- Top-down semi-join propagation from root to leaves in some topological order

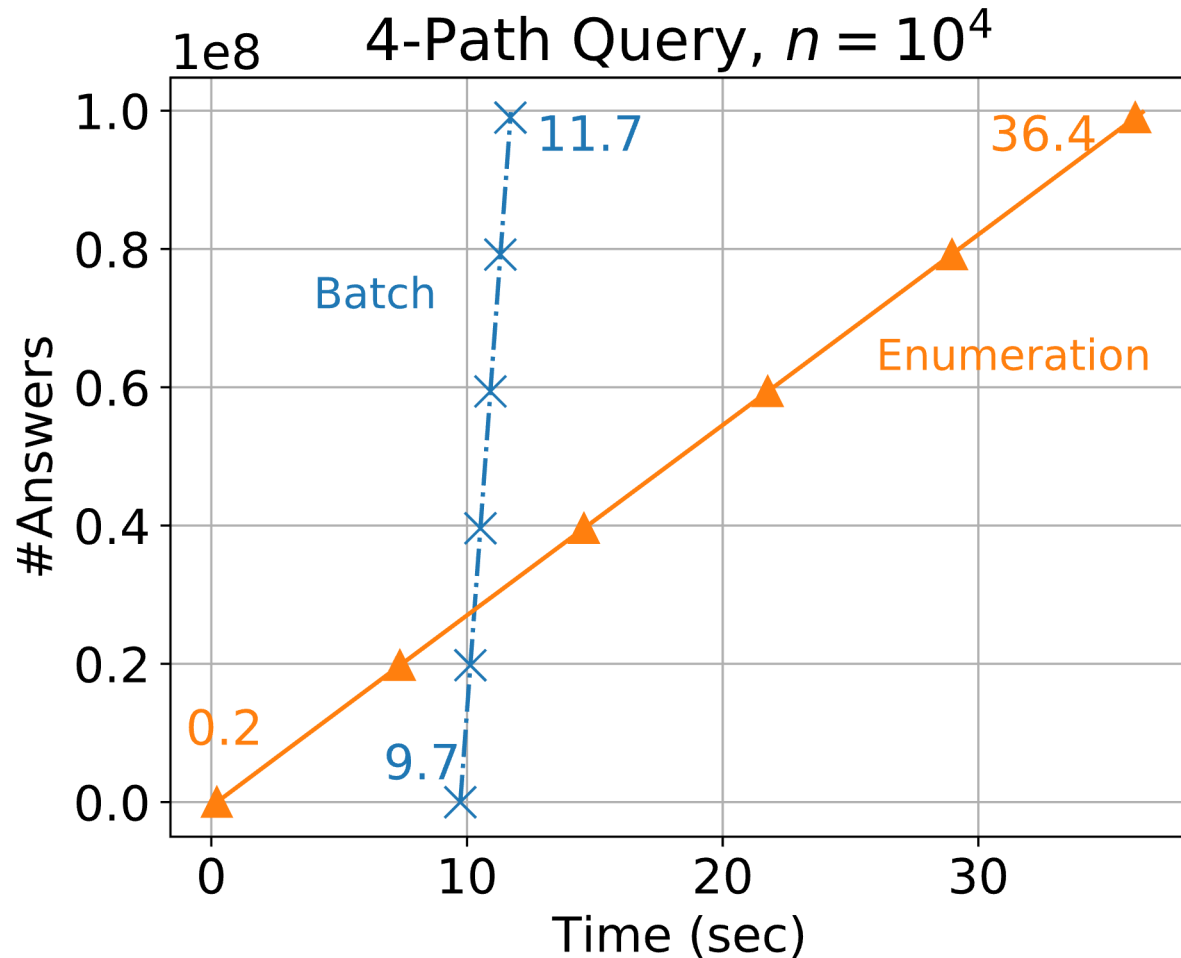
2. Enumeration phase \bowtie (compute answers with $O(1)$ delay)

- **Compute one result after the other in lexicographic order of the variables (added with the tables ordered in some topological order): $\{x,y\}+\{z,v\}+\{p\}+\emptyset+\{u\}$**

We start with some tuple in the root and extend it with consistent tuples from each table



Modified Yannakakis for output enumeration



- Standard Yannakakis:
 - Table-at-a-time / Breadth-first
 - After the semi-join reduction, Yannakakis visits each table once top-down, and at each stage increases the size of the answer set
- Enumeration:
 - Tuple-at-a-time / Depth-first
 - By a slight modification of Yannakakis and tuple by tuple

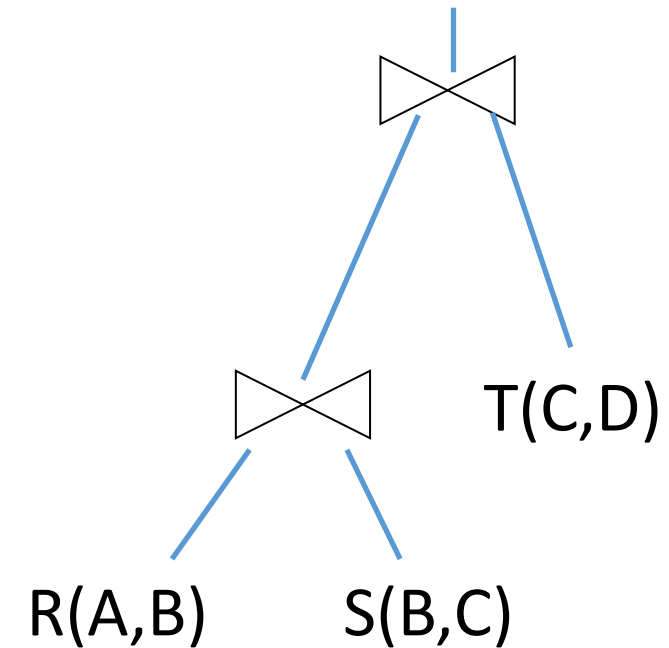
From Enumeration to Ranked Enumeration

Assume there is a preferred order on the answers

Q_3^∞ 3-chain query

R(A,B) S(B,C) T(C,D)

```
SELECT *  
FROM R natural join S  
natural join T
```

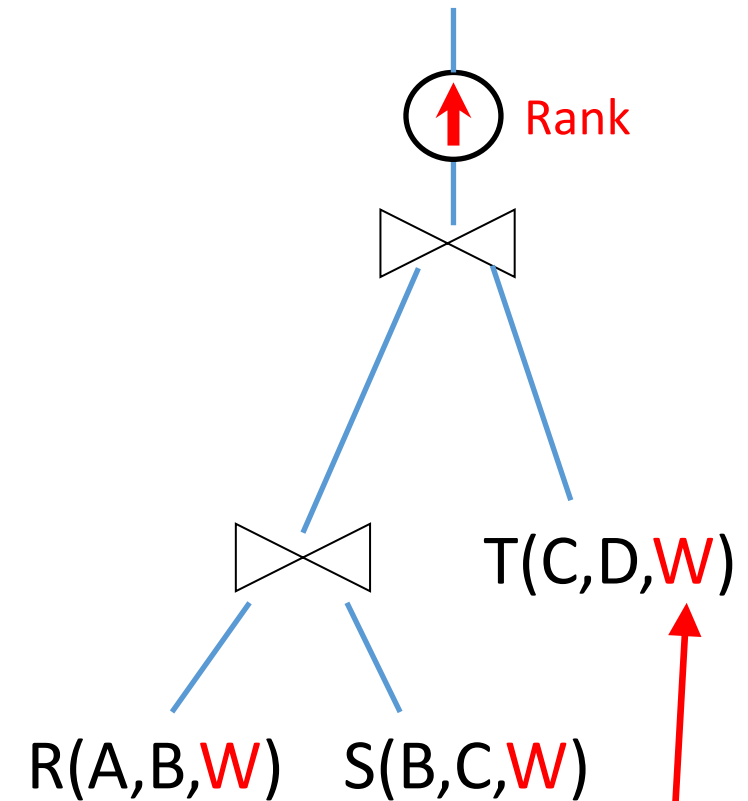


Assume there is a preferred order on the answers

Q_3^∞ 3-chain query

$R(A, B, W)$ $S(B, C, W)$ $T(C, D, W)$

```
SELECT *  
FROM R natural join S  
natural join T  
Order by R.W+S.W+T.W
```



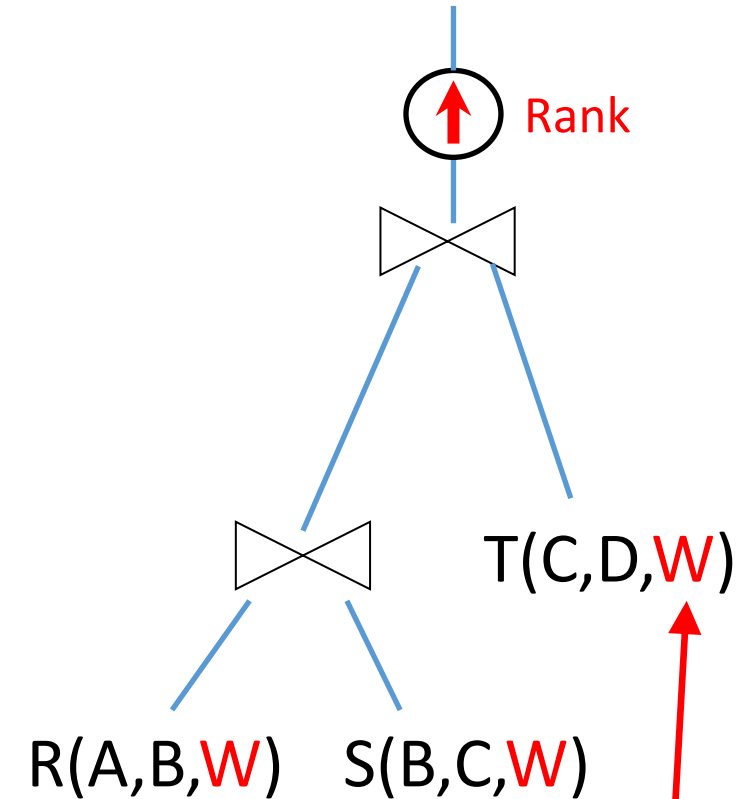
cost or weight associated with each tuple

Assume there is a preferred order on the answers

Q_3^∞ 3-chain query

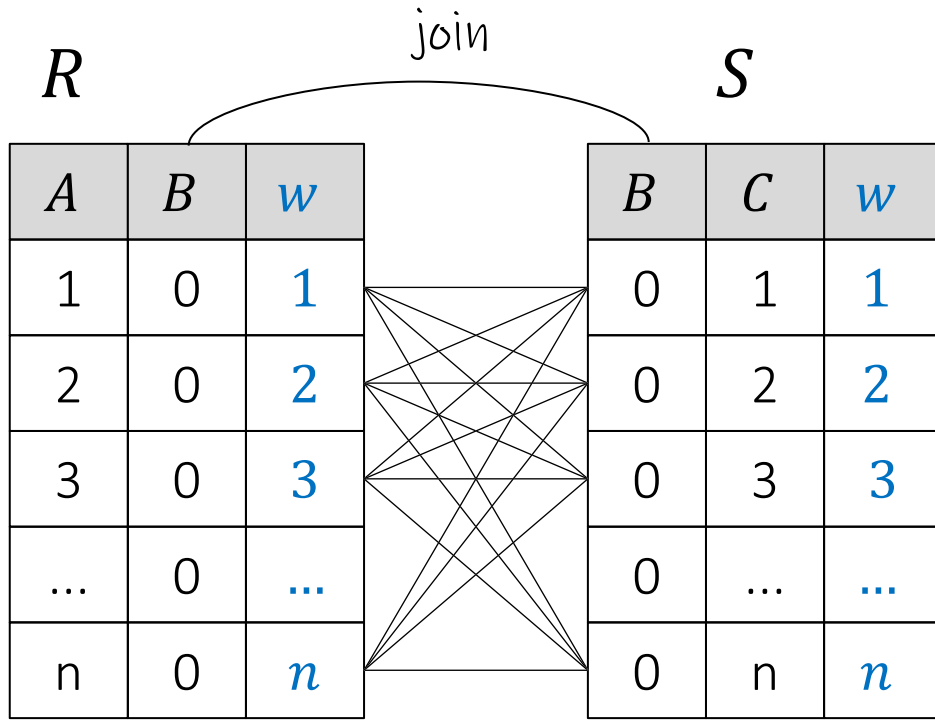
$R(A, B, W)$ $S(B, C, W)$ $T(C, D, W)$

```
SELECT *  
FROM R natural join S  
natural join T  
Order by R.W+S.W+T.W  
Limit 10
```



cost or weight associated with each tuple

Top- k is evaluated inefficiently by modern DBMS's



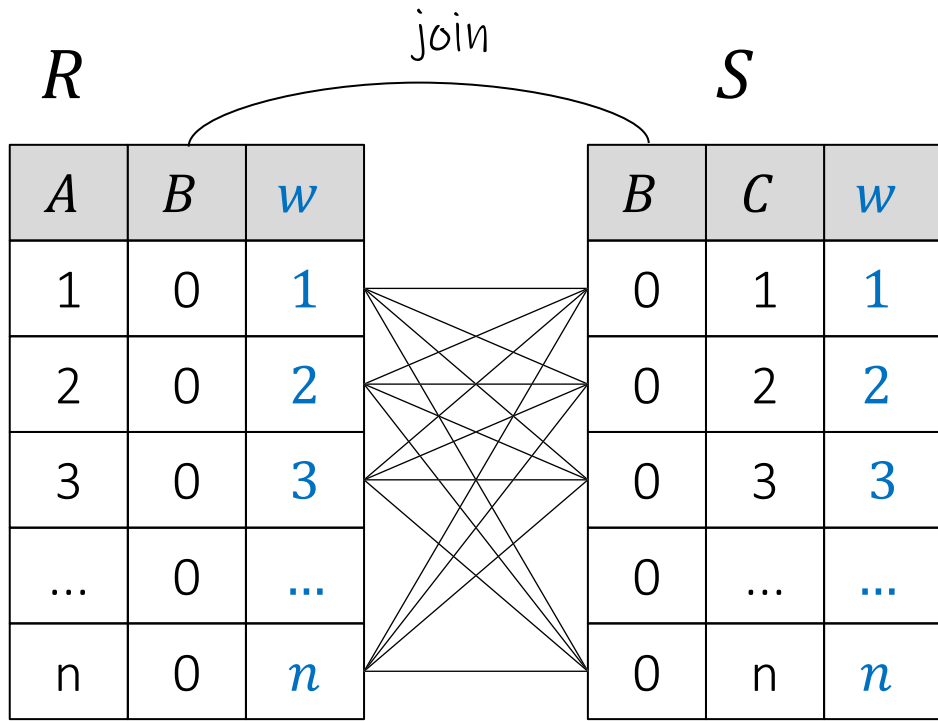
-- Query 1

```
SELECT  A, R.B, S.C,  
        R.W + S.W as weight  
FROM    R, S  
WHERE   R.B=S.B  
ORDER BY weight ASC  
LIMIT  1;
```

$n = 1,000:$ $t_{Q1} = 0.22 \text{ sec}$

$n = 10,000:$ $t_{Q1} = 22 \text{ sec}$

Top- k is evaluated inefficiently by modern DBMS's



Maximal intermediate result size is $O(n)$ 😊

~Dynamic programming

--- Query 1

```
SELECT  A, R.B, S.C,
        R.W + S.W as weight
FROM    R, S
WHERE   R.B=S.B
ORDER BY weight ASC
LIMIT  1;
```

$n=1,000:$ $t_{Q1}=0.22\text{ sec}$

$n=10,000:$ $t_{Q1}=22\text{ sec}$

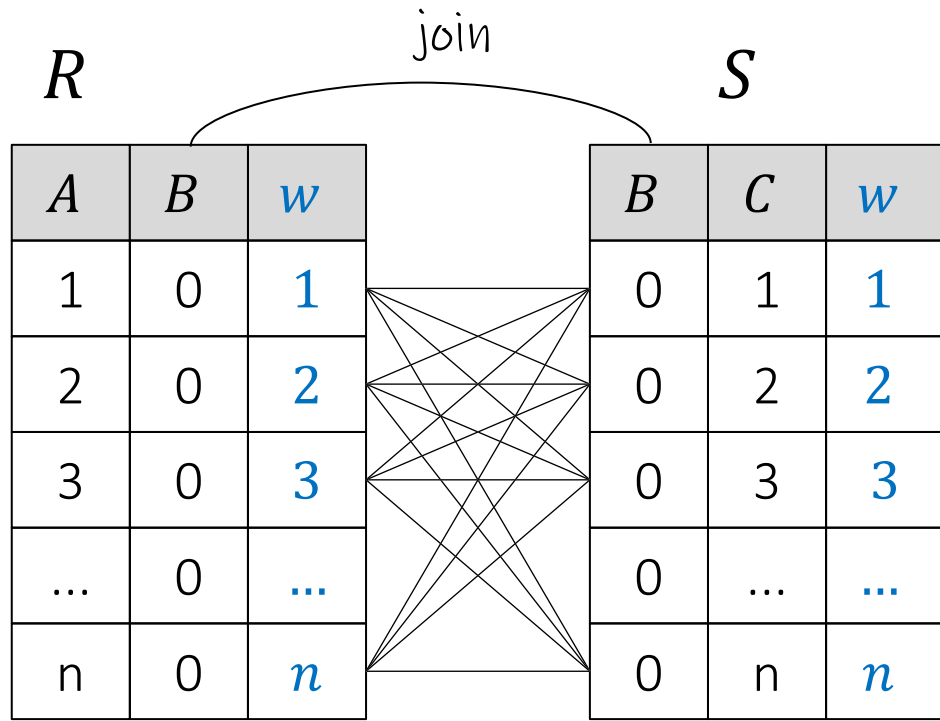
--- Query 2

```
SELECT R.A, X.B, S.C, X.W as weight
FROM R, S,
      (SELECT T1.B, W1, W2, W1+W2 W
       FROM
         (SELECT B, MIN(W) W1
          FROM R
          GROUP BY B) T1,
         (SELECT B, MIN(W) W2
          FROM S
          GROUP BY B) T2
       WHERE T1.B = T2.B
       ORDER BY W ASC
       LIMIT 1) X
WHERE X.B = R.B
AND X.W1 = R.W
AND X.B = S.B
AND X.W2 = S.W
LIMIT 1;
```

$t_{Q2}=1\text{ msec}$

$t_{Q2}=4\text{ msec}$

... even for just finding the minimum!



Maximal intermediate result size is $O(n)$ 😊

~Dynamic programming

-- Query 1

```
SELECT min(R.W + S.W) as weight
INTO record1
FROM R, S
WHERE R.B=S.B;
```

-- Query 2

```
SELECT min(W1+W2) as weight
INTO record2
FROM
  (SELECT B, MIN(W) W1
   FROM R
   GROUP BY B) T1,
  (SELECT B, MIN(W) W2
   FROM S
   GROUP BY B) T2
WHERE T1.B = T2.B;
```

n= 1,000: t_{Q1} = 0.1 sec

t_{Q2} <1 msec

n=10,000: t_{Q1} = 9.4 sec

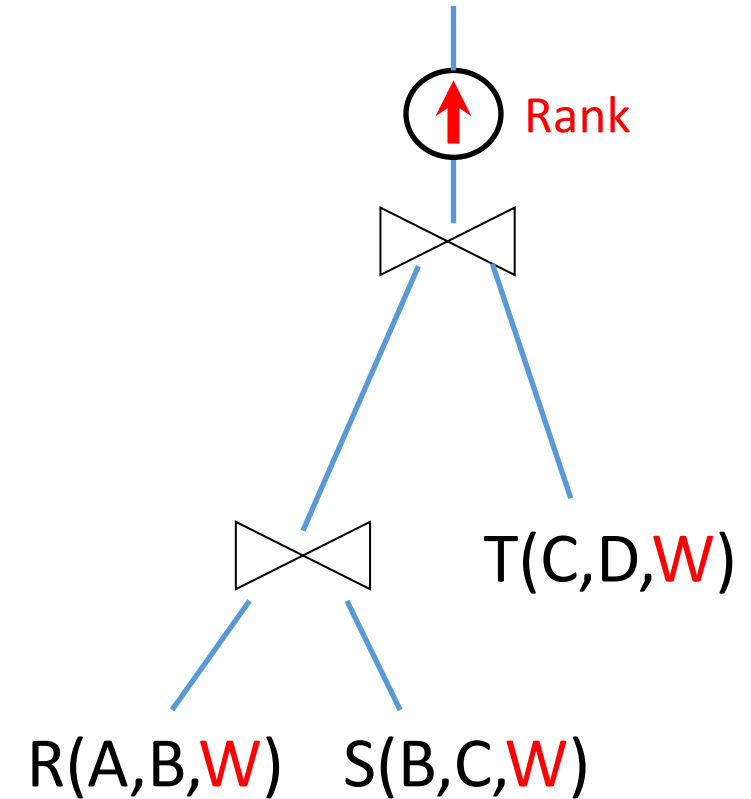
t_{Q2} =3 msec

Any- k : Faster and more versatile than Top- k

Q_3^∞ 3-chain query

$R(A, B, W)$ $S(B, C, W)$ $T(C, D, W)$

```
SELECT *  
FROM R natural join S  
natural join T  
Order by R.W+S.W+T.W  
Limit 10
```



Any- k : Faster and more versatile than Top- k

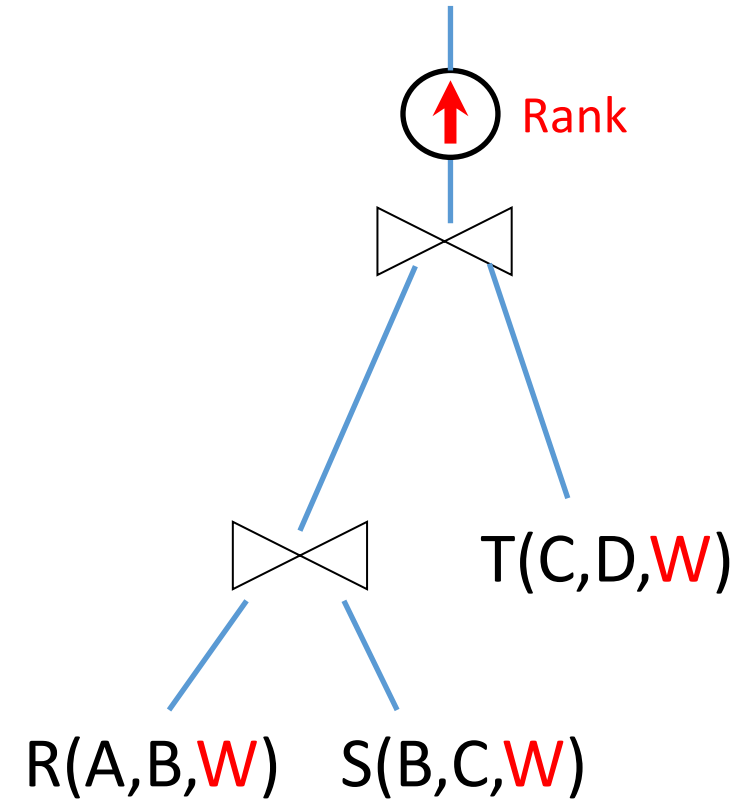
Q_3^∞ 3-chain query

$R(A, B, W)$ $S(B, C, W)$ $T(C, D, W)$

```
SELECT *  
FROM R natural join S  
natural join T  
Order by R.W+S.W+T.W  
Limit 10
```

Goal:

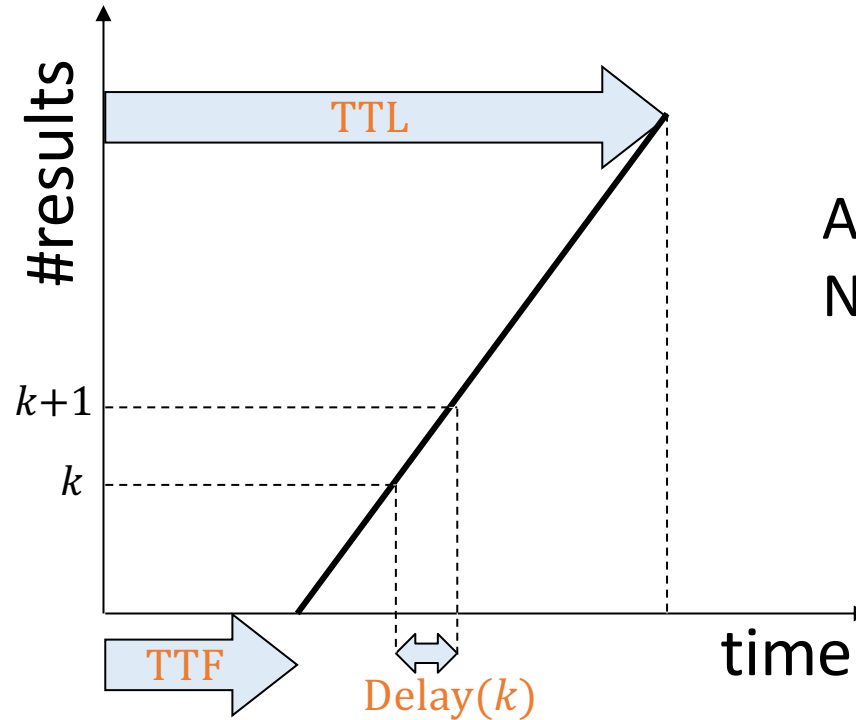
- Return the first result as fast as you can.
- Then the next.
- Then the next, ...
- Until the end.



Any- k (or "Ranked Enumeration"): Problem Definition

"Any- k ": Anytime algorithms + Top- k for Join Queries

Most important results first
(ranking function on output
tuples, e.g. sum of weights)



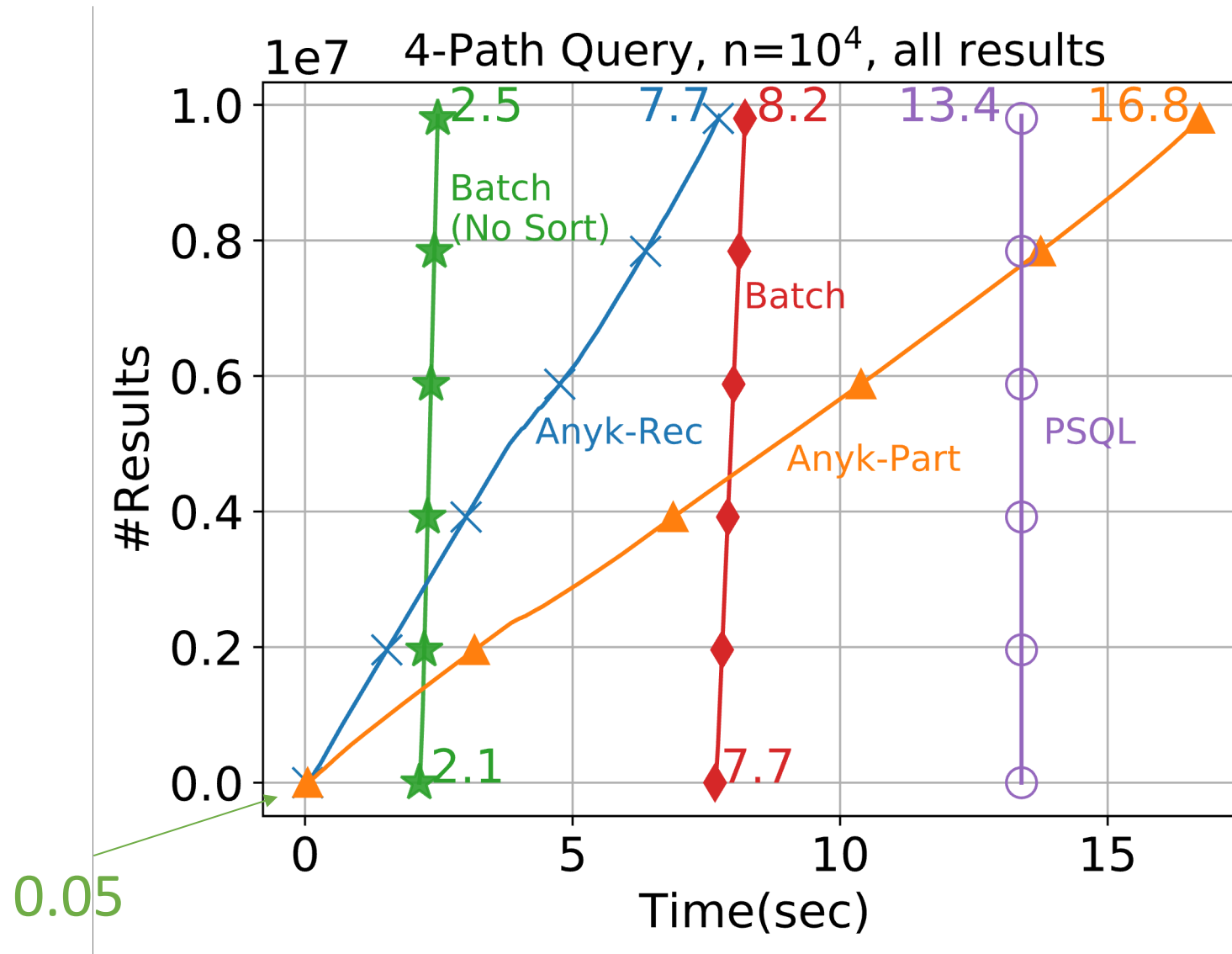
All results eventually returned
No need to set k in advance

RAM Cost Model:

- **TTF** = Time-to-First = $TT(1)$
- **Delay(k)** = Interval $k \rightarrow (k+1)$
- **TTL** = Time-to-Last = $TT(|out|)$

$TT(k)$ = Time-to- k^{th}

Experiments: TT(k) for Any- k variants vs. batch and PSQL

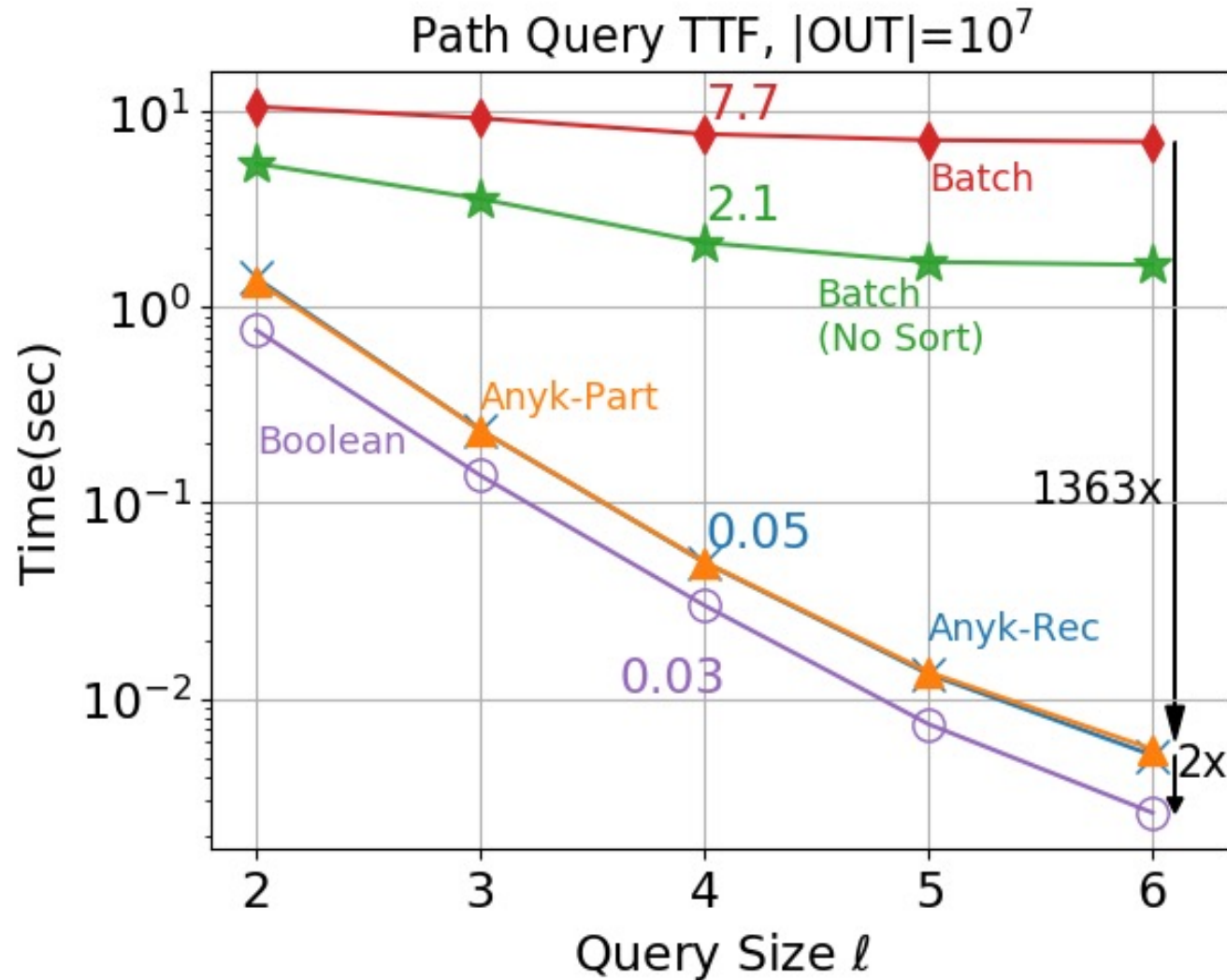


Source: <https://northeastern-datalab.github.io/anyk/>

Tziavelis, Ajwani, Gatterbauer, Riedewald, Yang. "Optimal Algorithms for Ranked Enumeration of Answers to Full Conjunctive Queries." PVLDB 2020. <https://doi.org/10.14778/3397230.3397250>

Wolfgang Gatterbauer. Principles of scalable data management: <https://northeastern-datalab.github.io/cs7240/>

Any- k : Faster and more versatile than Top- k



Path query with constant size output and increasing query size

How to deal with projections
in enumeration
("free-connex")

For α -acyclic queries, what changes for projections?

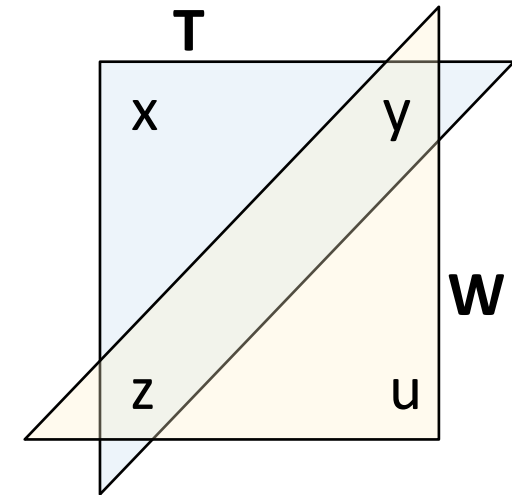
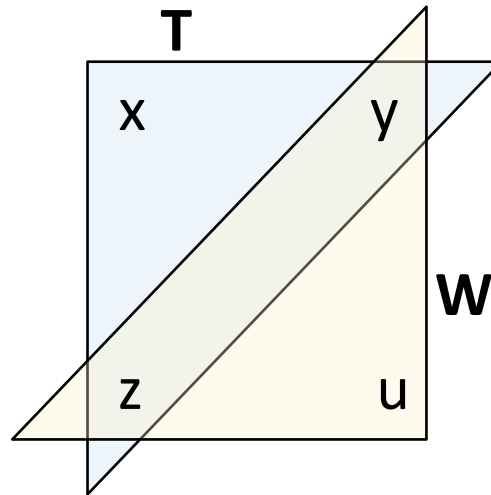
usually written as $O(n \cdot r)$ assuming output size $r \geq 1$

- Yannakakis works in $O(n + n \cdot r)$ for arbitrary projections
 - Then Enumeration works with linear delay and TTL is also $O(n + n \cdot r)$
- Yannakakis works in $O(n + r)$ for queries that are "free-connex"
 - Then Enumeration works with constant delay and TTL is also $O(n + r)$

$Q(x,y,z,u) :- T(x,y,z), W(u,y,z).$

$Q(x,y,z,u) :- T(x,y,z), W(u,y,z).$

Free-connex: the query remains *acyclic* after adding an edge for the set of projected variables



For α -acyclic queries, what changes for projections?



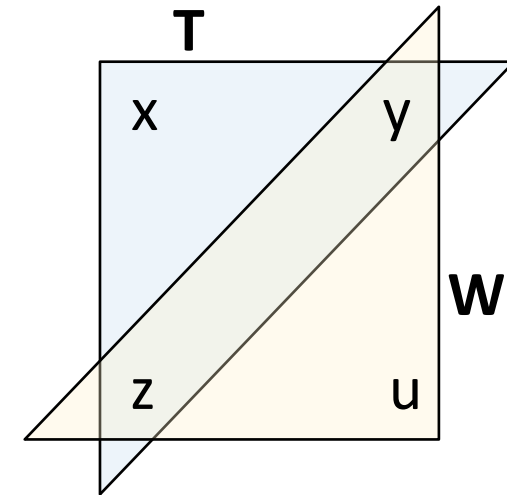
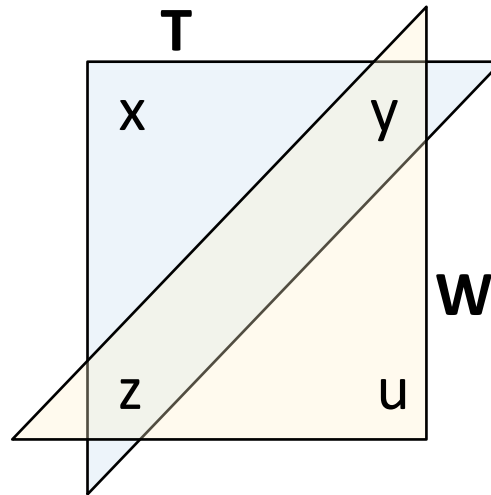
usually written as $O(n \cdot r)$ assuming output size $r \geq 1$

- Yannakakis works in $O(n + n \cdot r)$ for arbitrary projections
 - Then Enumeration works with linear delay and TTL is also $O(n + n \cdot r)$
- Yannakakis works in $O(n + r)$ for queries that are "free-connex"
 - Then Enumeration works with constant delay and TTL is also $O(n + r)$

$Q_1(x,y) :- T(x,y,z), W(u,y,z).$

$Q_2(x,u) :- T(x,y,z), W(u,y,z).$

Free-connex: the query remains *acyclic* after adding an edge for the set of projected variables



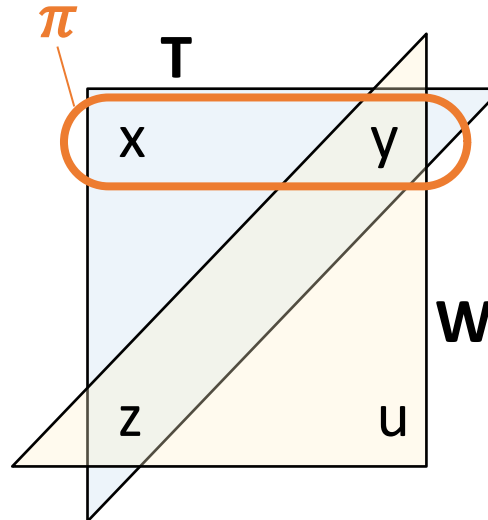
For α -acyclic queries, what changes for projections?



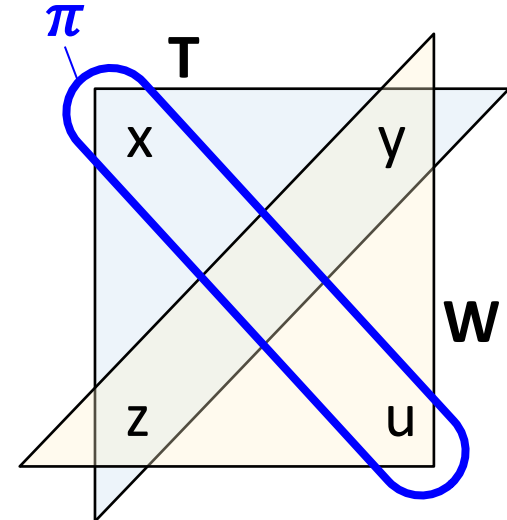
usually written as $O(n \cdot r)$ assuming output size $r \geq 1$

- Yannakakis works in $O(n + n \cdot r)$ for arbitrary projections
 - Then Enumeration works with linear delay and TTL is also $O(n + n \cdot r)$
- Yannakakis works in $O(n + r)$ for queries that are "free-connex"
 - Then Enumeration works with constant delay and TTL is also $O(n + r)$

$$Q_1(x, y) :- T(x, y, z), W(u, y, z).$$



$$Q_2(x, u) :- T(x, y, z), W(u, y, z).$$



Free-connex: the query remains acyclic after adding an edge for the set of projected variables

Which one is now free-connex ?

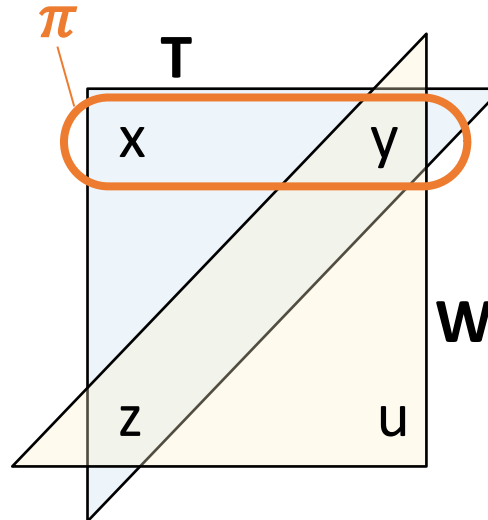
For α -acyclic queries, what changes for projections?

usually written as $O(n \cdot r)$ assuming output size $r \geq 1$

- Yannakakis works in $O(n + n \cdot r)$ for arbitrary projections
 - Then Enumeration works with linear delay and TTL is also $O(n + n \cdot r)$
- Yannakakis works in $O(n + r)$ for queries that are "free-connex"
 - Then Enumeration works with constant delay and TTL is also $O(n + r)$

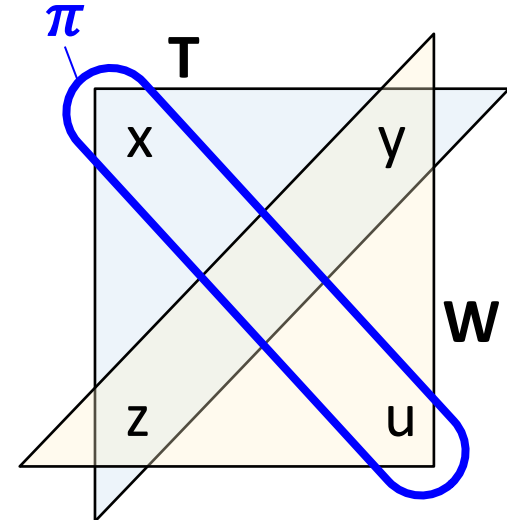
Free-connex: the query remains *acyclic* after adding an edge for the set of projected variables

$Q_1(x,y) :- T(x,y,z), W(u,y,z).$



free-connex 😊

$Q_2(x,u) :- T(x,y,z), W(u,y,z).$

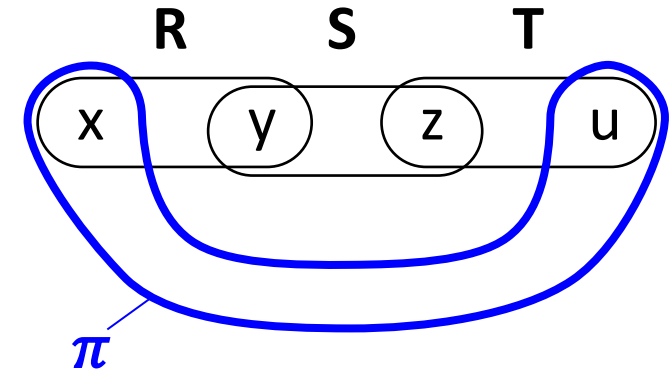
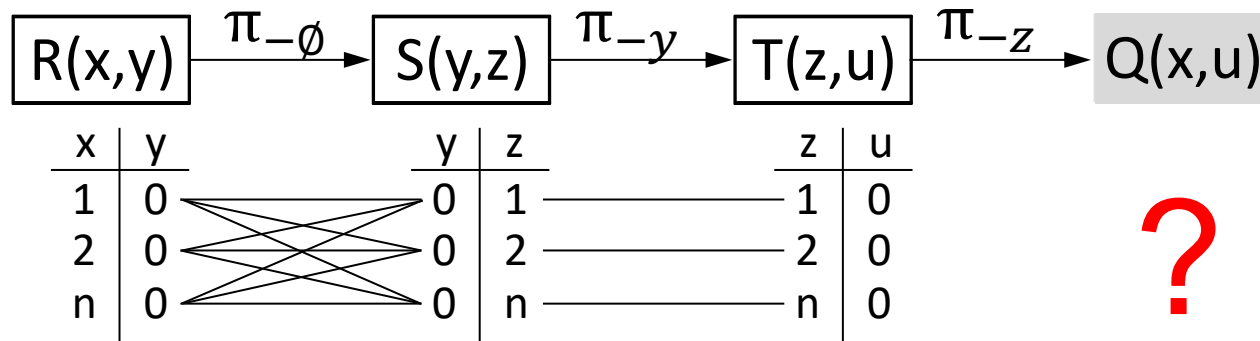


not free-connex ☹️

For α -acyclic queries, what changes for projections?

$Q(x,u) :- R(x,y), S(y,z), T(z,u).$

Join phase:

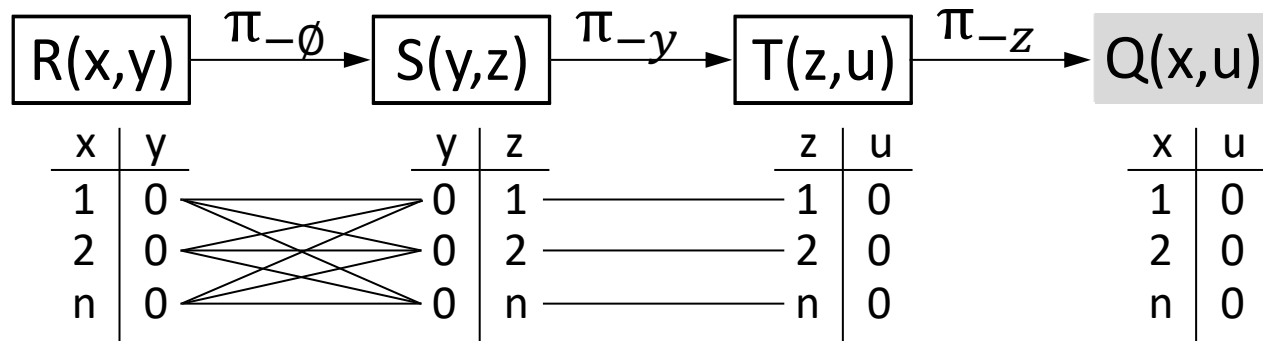


not free-connex ☹️

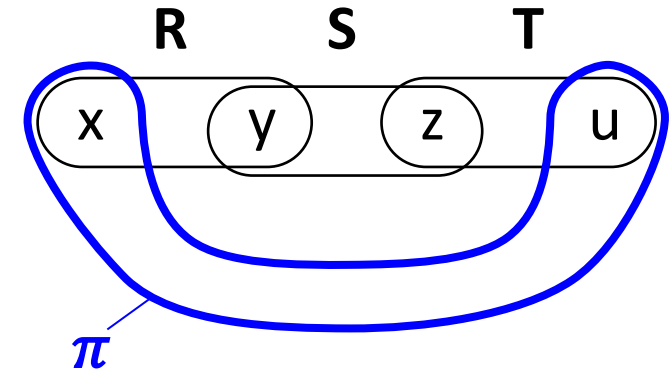
For α -acyclic queries, what changes for projections?

$Q(x,u) :- R(x,y), S(y,z), T(z,u).$

Join phase:



$O(n)$



not free-connex ☹️

$R(x,y) \bowtie S(y,z)$

x	y	z
1	0	1
2	0	1
n	0	1
...	0	...
n	0	3

$O(n^2)$

Output size is $r = O(n)$
 Join phase takes $O(n \cdot r) = O(n^2)$



Summary Acyclic queries

1. α (alpha)-acyclic queries imply a join tree can be found by GYO
2. Semi-join reductions work on trees. Can be extremely powerful and get rediscovered over and over again
3. Yannakakis works on acyclic queries and gives an optimal time guaranteed of $O(n+r)=O(|INPUT|+|Output|)$ for full CQs
 - $O(n+n \cdot r)$ for arbitrary projections
4. Enumeration slightly modifies Yannakakis to start returning results earlier (depth-first instead of breadth-first)