Databases

http://www.dcs.fmph.uniba.sk/~plachetk
/TEACHING/DB2

http://www.dcs.fmph.uniba.sk/~sturc/databazy/rldb

Tomáš Plachetka, Ján Šturc
Faculty of mathematics, physics and informatics
Comenius University, Bratislava
Summer 2023–2024

In this lecture, we will ignore negation in programs (to keep things simple). We will focus on **recursion** in programs (with functional symbols)

No difference in syntax, a **world of difference in computation** Examples:

- Simple join: q(X, Y, Z) ← r(X, Y), s(Y, Z). ?- q(a, b, c).
- Bottom-up (Datalog's naive evaluation) computes the join, then the selection:

$$B(X, Y, Z) = R(X, Y) \bowtie S(Y, Z); Q(X, Y, Z) = \sigma_{X=a \land Y=b \land Z=c} B(X, Y, Z)$$

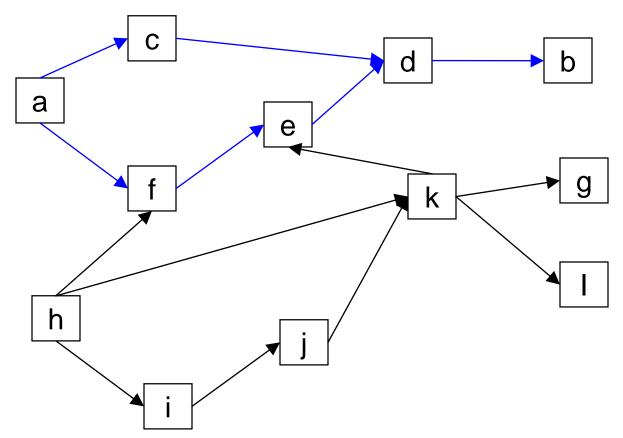
- Top-down (Prolog's SLD resolution) starts with the query. It attempts to prove q(a, b, c) by finding the rule whose head unifies with q(a, b, c) and finding instances of variables which satisfy all subgoals in the body of that rule
- · Recursion, a graph path:

$$p(X, Y) \leftarrow e(X, Y)$$
. $p(X, Y) \leftarrow e(X, Z)$, $p(Z, Y)$. ?- $p(a, b)$.

Top-down requires left recursion (EDB subgoals first), as it proves subgoals from left to right. The order is unimportant in bottom-up (conjunction/join is commutative). However, bottom-up computes all paths, although the query concerns only the path from a to b

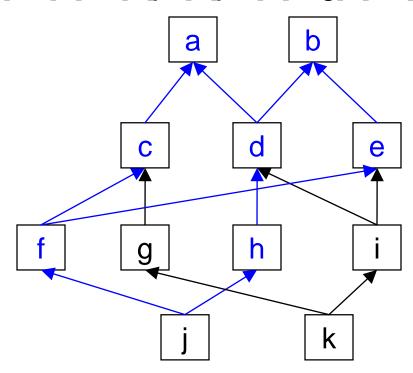
Example: recursion, a graph path $p(X, Y) \leftarrow e(X, Y). p(X, Y) \leftarrow e(X, Z), p(Z, Y).$?- p(a, b).

Bottom-up computation (Datalog) processes **all paths**, i.e. also paths in the black component of the graph. However, only blue edges matter



Another example: recursion, ancestors (an acyclic graph) anc(X, Y) \leftarrow par(X, Y). /* Y is parent of X */ anc(X, Y) \leftarrow par(X, Z), anc(Z, Y). **?- anc(j, A).** /* ancestors of j (j stands for John) */

EDB: par(X, Y) = {[c, a], [c, d], [d, b], [e, b], [f, c], [f, e], [g, c], [h, d], [i, d], [i, e], [j, f], [j, h], [k, g], [k, i]}



Addition of an arbitrary graph below vertices f, g, h, i, j, k does not influence the time complexity of top-down computation of John's ancestors. But it will influence the time complexity of bottom-up computation (naïve iteration computes the entire relation anc, not only John's ancestors)

Top-down computation: Rule-Goal Tree (RGT)

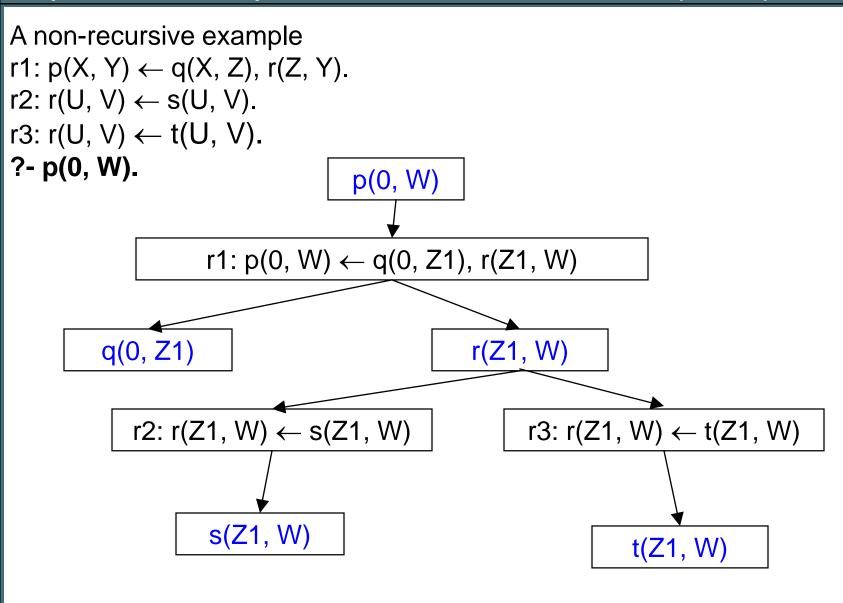
- 1. The goal, G_0 , is the root node of the tree.
- 2. The leaves of the root node are all rule nodes applicable to G_0 . Heads of these nodes are unified with G_0 so that:
 - a) Before the unification, all variables of the rules are made disjoint with variables appearing in the current goal (each rule gets "fresh" numbered variables before processing)
 - b) During unification with a head, goal variables are preferred (i.e. they do not disappear)
 - c) If a variable in the head is substituted, the substitution is applied to all occurrences of that variable in the rule
 - d) Variables not appearing in the head (the fresh variables) are local in the rule (i.e. they do not appear in any other rule)
- 3. The leaves of a rule node are subgoals of the rule body. Each subgoal is recursively processed in a similar manner as G₀

. . .

Top-down computation: Rule-Goal Tree (RGT)

```
Example: recursion, ancestors
r1: anc(X, Y) \leftarrow par(X, Y). /* Y is parent of X */
r2: anc(X, Y) \leftarrow par(X, Z), anc(Z, Y).
?- anc(j, A). /* ancestors of j (j stands e.g. for John) */
                                  anc(j, A)
      r1: anc(j, A) \leftarrow par(j, A)
                                               r2: anc(j, A) \leftarrow par(j, Z1), anc(Z1, A)
                                             par(j, Z1)
                                                                    anc(Z1, A)
               par(j, A)
     r1: anc(Z1, A) \leftarrow par(Z1, A)
                                              r2: anc(Z1, A) \leftarrow par(Z1, Z2), anc(Z2, A)
               par(Z1, A)
                                              par(Z1, Z2)
                                                                     anc(Z2, A)
```

Top-down computation: Rule-Goal Tree (RGT)



Top-down: propagation of bindings and results in RGT

- Each **goal node** is assigned a **binding relation M** ("magic **predicate**"). Bound variables of the goal are attributes of the corresponding magic relation. The contents of the magic relation is a finite set of constant values resulting from the previous computation ("sideway-passed information")
- In each rule node, i-th argument of the rule is assigned a "supplementary relation" S_i. Supplementary relations collect sideway-passed information, i.e. possible instantiations for variables of the rule. Initially, this information is determined only by the head of the rule. It is updated as the computation moves from left to right over subgoals of the rule
- Each goal node as well as each rule node yield a resulting relation. For a goal node, it is the set of tuples which satisfy the goal. For a rule node, it is the set of tuples which satisfy the head of the rule

Top-down: actual computation

- Actual computation takes place in rule nodes
- In the root and inner **goal nodes**, the resulting relation is computed as **union of the results** from the child nodes
- Leaves of the tree are joins (selections) of the magic relations with EDB relations
- In rule nodes, the subgoals are first converted to supplementary relations with variables (using atov). The supplementary relations are joined (or antijoined, in case of negation). The result is then converted to head (using vtoa) and sent to the parent node

Top-down: RGT construction and evaluation

RGT construction and evaluation

- Input: a set of safe rules, EDB and a query (goal) G₀
- Output: a set of all tuples which satisfy G₀
- Method: two mutually recursive functions, expand_goal and expand_rule
 - expand_goal(M, G, R), where G is a goal, M is the binding relation assigned to G, R is the resulting relation
 - expand_rule(S_0 , r, R), where r is a rule, S_0 is the supplementary predicate containing bindings given by the head, R is the resulting relation

For a query ?- G_0 , the computation begins with expand_goal(M_0 , atov(G, P), R), where M_0 is the binding relation for the variables in G_0 , P is the predicate in G_0 , R is the result. (For a query with no bindings, M_0 is initialised to a universe, i.e. $S \bowtie M_0 = S$ for an arbitrary relation S.)

Top-down: expand_goal(M, G, R)

```
expand_goal(M, G, R) {
    if (G is a goal with an EDB predicate P)
         R = M \bowtie atov(G, P);
    else {
        /* G is a goal with an IDB predicate */
         R = \emptyset; /* the result will be accumulated in R */
         for (each rule r whose head H unifies with G) {
             \tau = mgu(G, H);
             H' = \Pi_M (H\tau); /* H\tau converted to arguments of G */
             S_0 = atov(H', M);
             expand_rule(S_0, r\tau, R_r);
             R = R \cup R_r;
```

Top-down: expand_rule(S₀, r, R)

```
expand_rule(S_0, r, R) {
     /* let r be of form H \leftarrow G_1, ..., G_k */
     for (i = 1; i \le k; i++)
          M_i = vtoa(G_i, S_{i-1});
           expand_goal(M<sub>i</sub>, G<sub>i</sub>, R);
           Q_i = atov(G_i, R); /* convert R_i to variables */
          S_i = \prod_T (S_{i-1} \bowtie Q_i); /* T is the set of variables which
                appear in S<sub>i-1</sub> or Q<sub>i</sub> in the rule r */
     R = vtoa(H, S_k);
```

Top-down: QRGT algorithm

- Recursive expand_rule / expand_goal algorithm is a depth-first traversal of the RGT. It can end in an infinite loop
- Modification of the expand_rule / expand_goal algorithm:
- QRGT (queue-based rule-goal-tree expansion). The idea is that breadth-first traversal of RGT does not get lost in an infinite branch
- QRGT computes the fix-point for safe Datalog programs
- For a goal G with magic relation M, QRGT computes a tuple $[t_1, ..., t_k]$ when this tuple is computed by the bottom-up computation

Adorned predicates and rules

- Adornment for a predicate is a (finite) string of symbols 'b' and 'f' which stand for 'bound' and 'free'. An adornment states which arguments of the predicate are bound just before the predicate is evaluated (called)
- Adornment for a rule (more precisely, for a position inside a rule) is a list assigned to "spaces between goals" in the rule,

 $[V_{b1}, V_{b2}, ...V_{bm} \mid V_{f1}, V_{f2}, ..., V_{fn}]$. The symbol '|' in the list separates bound and free variables of the rule when the computation just after the top-down computation has reached that point inside the rule. Note that a bound variable inside a rule is restricted to finitely many values. A variable is bound after the goal G_i when it has already been bound in the head of the rule, or when it appears in the rule anywhere before the goal G_i (from left to right, including G_i)

Rule-Goal Graph consists of goal (predicate) nodes and rule position nodes.

RGG is a generalisation of RGT. Unlike RGT, RGG is always finite, but may contain cycles (when a predicate or a rule goal with the same adornment is evaluated more than once during the top-down computation). Edges of RGG connect nodes so that:

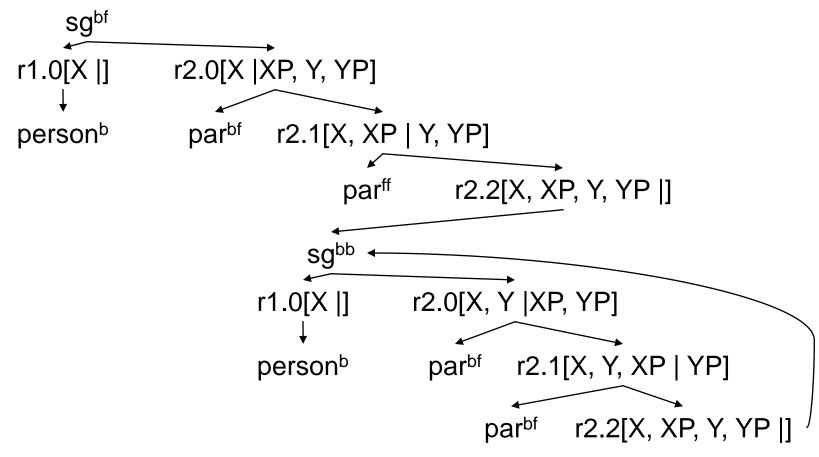
- Node with an EDB predicate has no outgoing edges
- Outgoing edges of a node with an adorned IDB predicate p^a end in rule nodes r₀[... | ...] such that the head of the rule can be unified with p^a, i.e. they bind the same variables
- Outgoing edges of a node with an adorned rule position $r_i[\dots \mid \dots]$ end in
 - a) goal nodes p_i^a , where p_i is the predicate in goal G_i (i.e. the goal following the position r_i) and the sets of variables bound in a and in the adornment of r_i are equal
 - b) node r_{i+1} (if r_i is a rule position before the penultimate goal), where variables bound in r_{i+1} are those which have been bound in r_i plus those which appear in the goal G_i

Example: the same generation (person and parent are EDB predicates)

```
r1: sg(X, X) \leftarrow person(X).
```

r2: $sg(X, Y) \leftarrow par(X, XP)$, par(Y, YP), sg(XP, YP).

?- sg(j, Y).

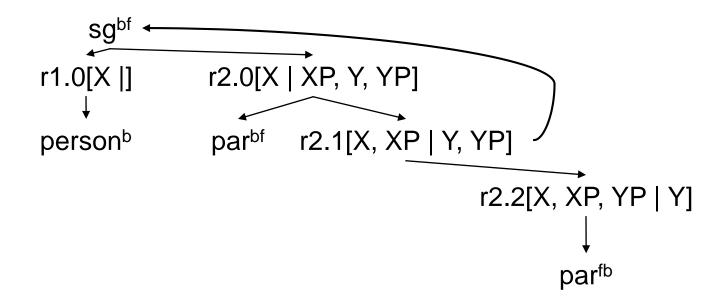


Example: the same generation with reordered goals:

r1: $sg(X, X) \leftarrow person(X)$.

r2: $sg(X, Y) \leftarrow par(X, XP)$, sg(XP, YP), par(Y, YP).

?-sg(j, Y).



Problems arising during the construction of an RGG:

- Rectification of subgoals in rules. This is advisable when either a constant or repeated variable appears in an IDB goal. (For example, adornment for p(X, X) is neither pff, nor pbb, it is actually slightly stronger. Similarly, p(1, 3) is stronger than just pbb.)
- Making adornments of IDB predicates uniform by reordering subgoals in rules
- Making built-in subgoals feasible. It may be possible to compute built-in subgoals for some adornments, but not all
- → feasibility problem for RGG

Making adornments of IDB predicates uniform

- For each adorned predicate p^{α} in RGG, create a new predicate p_α
- Copy all rules r with head p into rules r _α with head p_α
- In all goal nodes r_0 , ..., r_k of the transformed rule r_0 , modify its child nodes as follows:
 - a) Keep EDB and built-in predicates in the rule r $_\alpha$ as they were in rule r
 - b) Change all adorned IDB predicates q^β to q_β

Making adornments of IDB predicates uniform: example

Original program:

- r1: $sg(X, X) \leftarrow person(X)$.
- r2: $sg(X, Y) \leftarrow par(X, XP)$, par(Y, YP), sg(XP, YP).
- ?- sg(j, Y).

Modified program with uniform adornments of IDB predicate sg:

- r1_bf: $sg_bf(X, X) \leftarrow person(X)$.
- r2_bf: sg_bf(X, Y) \leftarrow par(X, XP), par(Y, YP), sg_bb(XP, YP).
- r1_bb: $sg_b(X, X) \leftarrow person(X)$.
- r2_bb: $sg_b(X, Y) \leftarrow par(X, XP)$, par(Y, YP), $sg_b(XP, YP)$.
- ?- sg_bf(j, Y).

Rectification

Motivation:

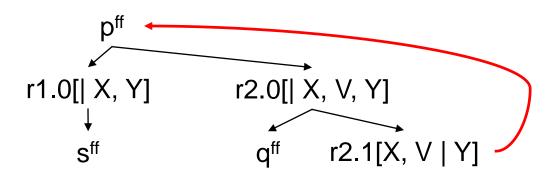
r1: $p(X, Y) \leftarrow s(X, Y)$.

r2: $p(X, Y) \leftarrow q(X, V), p(Y, Y)$.

?-p(X, Y).

Note that the adornment pff does not fully capture the bindings in the call p(Y, Y) in r2. Although the variable Y is free, the two arguments in p are not arbitrary, they must be equal!

The adornments in the following RGG are not quite exact



Rectification

Rectification is an algorithm which produces a program without duplicated variables and without constants in IDB subgoals.

It defines **new predicates** (with fewer arguments):

```
r1: p(X, Y) \leftarrow s(X, Y).
```

r2:
$$p(X, Y) \leftarrow q(X, V), p(Y, Y)$$
.

$$p(X, Y) \leftarrow s(X, Y)$$
.

$$p(X, Y) \leftarrow q(X, V), p1(Y).$$
 /* $p1(Y) \leftarrow p(Y, Y)$ */

$$p1(Y) \leftarrow s(Y, Y)$$
. /* expansion using r1 */

$$p1(Y) \leftarrow q(Y, V), p1(Y)$$
. /* expansion using r2 */

Rectification

```
while (IDB goal g exists with a duplicated variable or a constant in
arguments)
   Let g be a goal where a predicate p is called. Replace p in the
     body a new predicate p' with no duplicated variables and no
     constants in arguments
   For each rule r with head h, create a new rule r'. To do that,
     compute \tau = mgu(g, h), preferring the variables of g. The
     new rule r' is r\tau, where all occurences of p are replaced with
     p'.
   In all the other rules, replace p with p'.
```

Note that this algorithm always terminates (the number of arguments decreases in each iteration). Moreover, if the original program is safe, so is the rectified program

A rule in RGG is evaluated always from left to right. However, we can arbitrarily order the goals in the rule

Sometimes we must reorder the goals because of built-in goals

For example, the rule

$$p(X, Y) \leftarrow X < Y, r(X, Y).$$

equivalently: $p(X, Y) \leftarrow less(X, Y), r(X, Y)$.

cannot be evaluated from left to right e.g. for the adornment pff.

The only adornment allowed in a call to less is lessbb.

But this ordering of goals is all right:

$$p(X, Y) \leftarrow r(X, Y), less(X, Y).$$

Other reasons for reordering goals:

- Making use of indexes for EDB. Some adornments of EDB predicates may profit from using an index attached to a database relation
- IDB predicates with functional symbols in their definition may also forbid some adornments. For example, nat^f leads to an infinite expansion, whereas nat^b does not.
- Rules for nat(.): nat(0). nat(s(X)) \leftarrow nat(X).
- · Negated IDB goals require that all their arguments are bound

We assume that **bound** is easier

We define a partial ordering of adornments:

 $\alpha \leq \beta$ if β has 'b' at least on those positions where α has

Then if $\alpha \leq \beta$ and we can evaluate p^{α} , then we can also evaluate p^{β}

For example, if we can evaluate pf, then we can also evaluate pb

We say that an adornment is **allowed** if the predicate can be evaluated for that adornment

For each predicate, there is a set for minimal allowed

adornments

Assumed that for each adorned rule (i.e. adornment of the head), minimal allowed adornments for each goals are given

How to find an ordering of the goals so that the rule can be evaluated?

Backtracking: enumerate all the orderings and test whether a feasible

RGG (only with allowed adornments) can be constructed

Optimisations:

- Assume "bound is easier"
- Maintain two sets of adorned goals: 1.a set T of target adorned goals (this initially contains only the query); 2.a set F of adorned goals which cannot be realized (this initially contains goals with adornments which must be avoided, i.e. which are not allowed)

Example: the same generation

r1: $sg(X, X) \leftarrow person(X)$.

r2: $sg(X, Y) \leftarrow par(X, XP)$, par(Y, YP), sg(XP, YP).

?- sg(john, W).

T={sgbf}, F={parff, personf}

There is only one ordering for r1bb which avoids personf.

For r2bf, this ordering must be tested: par₁bf, sgfb, par₂fb. Therefore, sgfb

is added into the set T. It remains to show that sgfb can be realised.

There is only one ordering for r1bb which avoids personf.

For r2fb, the following ordering is realisable: par2bf, sgbf, par1fb.

We end up with T={sgbf, sgfb}. All goals in T are realisable

```
Example: the same generation, ?- sg(john, W).
                                                        F={parff, personf}
r1: sg(X, X) \leftarrow person(X).
r2^{bf}: sg(X, Y) \leftarrow par(X, XP), sg(XP, YP), par(Y, YP).
r2^{fb}: sg(X, Y) \leftarrow par(Y, YP), sg(XP, YP), par(X, XP).
                             sg<sup>bf</sup> ←
Feasible RGG:
                                       r2.0[X |XP, Y, YP]
       r1.0[X |]
                                parbf
                                                 r2.1[X, XP | Y, YP]
       personb
                                      sgfb
                                                          r2.2[X, XP, YP | Y]
                                                                  parfb
                        r1.0[X |]
                                        r2.0[Y | X, XP, YP]
                                     parbf
                                             r2.1[Y, YP | X, XP]
                        personb
                                                            r2.2[Y, YP, XP | X]
                                                                   parfb
```