On the problem of computing the well-founded semantics*

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Abstract

The well-founded semantics is one of the most widely studied and used semantics of logic programs with negation. In the case of finite propositional programs, it can be computed in polynomial time, more specifically, in $O(|At(P)| \times size(P))$ steps, where size(P) denotes the total number of occurrences of atoms in a logic program P. This bound is achieved by an algorithm introduced by Van Gelder and known as the alternating-fixpoint algorithm. Improving on the alternating-fixpoint algorithm turned out to be difficult. In this paper we study extensions and modifications of the alternating-fixpoint approach. We then restrict our attention to the class of programs whose rules have no more than one positive occurrence of an atom in their bodies. For programs in that class we propose a new implementation of the alternating-fixpoint method in which false atoms are computed in a top-down fashion. We show that our algorithm is faster than other known algorithms and that for a wide class of programs it is linear and so, asymptotically optimal.

KEYWORDS: well-founded semantics, alternating fixpoint

1 Introduction

The well-founded semantics was introduced (Van Gelder *et al.*, 1991) to provide 3-valued interpretations to logic programs with negation. Since its introduction, the well-founded semantics has become one of the most widely studied and most commonly accepted approaches to negation in logic programming (Alferes *et al.*, 1995; Fitting, 1991; Chen *et al.*, 1995; Chen & Warren, 1996; Zukowski *et al.*, 1997; Brass & Dix, 1998). It was implemented in several top-down reasoning systems, most prominent of which is XSB (Rao *et al.*, 1997).

The well-founded semantics is closely related to the stable-model semantics (Gelfond & Lifschitz, 1988), another major approach to logic programs with negation. The well-founded semantics approximates the stable-model semantics (Van Gelder *et al.*, 1991; Fitting, 2001). Moreover, computing the well-founded model of propositional programs is polynomial (Van Gelder, 1989) while computing stable models is NP-hard (Marek & Truszczyński, 1991). Consequently, evaluating the well-founded

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semantics can be used as an effective preprocessing technique in algorithms to compute stable models (Subrahmanian *et al.*, 1995). In addition, as demonstrated by smodels (Niemelä & Simons, 1996), at present the most advanced and most efficient system to compute stable models of DATALOG¬ programs, the well-founded semantics can be used as a powerful lookahead mechanism.

Despite the importance of the well-founded semantics, the question of how fast it can be computed has not attracted significant attention. Van Gelder (1989) described the so called *alternating-fixpoint* algorithm. Van Gelder's algorithm runs in time $O(|At(P)| \times size(P))$, where At(P) is the set of atoms occurring in a logic program P, |At(P)| denotes the cardinality of At(P), and size(P) is the size of P (the total number of atom occurrences in P). Improving on this algorithm turned out to be difficult. The first progress was obtained by Berman *et al.* (1995). The algorithm described there, when restricted to programs whose rules contain at most two positive occurrences of atoms in their bodies, runs in time $O(|At(P)|^{4/3}|P|^{2/3} + size(P))$, where |P| is the number of rules in P. For programs whose rules have no more than one positive atom in the body a better estimate of $O(|At(P)|^{3/2}|P|^{1/2} + size(P))$ was obtained. For some classes of programs this is an asymptotically better estimate than the $O(|At(P)| \times size(P))$ estimate that holds for the algorithm by Van Gelder.

A different approach to computing the well-founded model has also been proposed (Zukowski et al., 1997; Brass et al., 2001). It is based on the notion of a program transformation (Brass & Dix, 1998). The authors describe there several transformations that can be implemented in linear time and that simplify a program while (essentially) preserving the well-founded semantics. These transformations are: the positive reduction, success, negative reduction and failure (PSNF transformations, for short). They allow one to compute in linear time the Kripke-Kleene semantics (Fitting, 1985) of the program. To compute the well-founded semantics one also needs to detect the so-called positive loops. The complexity of this task dominates the asymptotic complexity of the well-founded semantics computation. No improved algorithms for the positive-loop detection are offered in Brass et al. (2001) so the worst-case asymptotic complexity of the algorithm presented there remains the same as that of the alternating-fixpoint method. However, due to the use of PSNF transformations, that simplify the program, the algorithm based on program transformations may in practice run faster. In contrast to the approach studied in Brass et al. (2001), we focus here on the positive-loop detection task.

The alternating-fixpoint algorithm works by successively improving lower approximations T and F to the sets of atoms that are true and false (under the well-founded semantics), respectively. The algorithm starts with $T = \emptyset$. Using this estimate, it computes the first estimate for F. Next, using this estimate, in turn, it computes a better estimate for T. The algorithm continues until further improvements are not possible. It returns the final sets T and F as the well-founded semantics. A dual version of the alternating-fixpoint algorithm, starting with $F = \emptyset$ and then alternatingly computing approximations to T and F, is also possible. The most time-consuming part of this algorithm is in computing estimates to the set of atoms that are false (in this part, in particular, positive loops are detected). In the Van Gelder algorithm, the best

possible approximation (given the current estimate for T) is always computed by using a bottom-up approach.

In this paper, we focus on the problem of detecting positive loops and computing new false atoms. We restrict our attention to the class of programs that have at most one positive atom in the body. We denote this class of programs by \mathcal{LP}_1 . We show that for programs from \mathcal{LP}_1 , false atoms can be computed by means of a top-down approach by finding atoms that do not have a proof. Moreover, we show that it is not necessary to find all atoms that can be established to be false at a given stage. Finding a proper subset (as long as it is not empty) is also sufficient and results in a correct algorithm. We apply these techniques to design a version of an alternating-fixpoint algorithm computing the well-founded semantics of programs from the class \mathcal{LP}_1 . We demonstrate that the resulting algorithm is asymptotically better than the original alternating-fixpoint algorithm by Van Gelder. Specifically, we show that our algorithm runs in time $O(|At(P)|^2 + size(P))$. Thus, for programs with $size(P) \ge |At(P)|^2$, our algorithm runs in linear time and is asymptotically optimal! It is also easy to see that when |P| > |At(P)|, the asymptotic estimate of the running time of our algorithm is better than that of algorithms by Van Gelder (1989) and Berman et al. (1995).

As mentioned above, our approach is restricted to the class \mathcal{LP}_1 . Applicability of our method can, however, be slightly extended. Let us denote by \mathcal{LP}_1^+ the class of these logic programs that, after simplifying by means of PSNF transformations (or, equivalently, with respect to the Kripke–Kleene semantics) fall into the class \mathcal{LP}_1 . Since PSNF transformations (the Kripke–Kleene semantics) can be computed in linear time, the asymptotic estimate of the running time of our method extends to all programs in the class \mathcal{LP}_1^+ .

The paper is organized as follows. In the next section we provide a brief review of the key notions and terminology. In section 3 we describe several modifications to the original Van Gelder algorithm, we show their correctness and estimate their running time. The ultimate effect of our considerations there is a general template for an algorithm to compute the well-founded semantics. Any algorithm computing some (not necessarily all) atoms that can be established as false given a current estimate to the well-founded can be used with it. One such algorithm, for programs from the class \mathcal{LP}_1 , is described and analyzed in section 4. It constitutes the main contribution of the paper and yields a new, currently asymptotically most efficient algorithm for computing the well-founded semantics for programs in \mathcal{LP}_1 . The last section contains conclusions.

2 Preliminaries

We start by reviewing basic concepts and notation related to logic programs and the well-founded semantics, as well as some simple auxiliary results. In the paper we consider the propositional case only.

Let P be a normal logic program. By At(P) we denote the set of atoms occurring in P. Let $M \subseteq At(P)$ (throughout the paper we often drop a reference to P from our notation, whenever there is no danger of ambiguity). By P_M we denote the program

obtained from P by removing all rules whose bodies contain negated literals of the form not(a), where $a \in M$. Further, by P^h we denote the program obtained from P by removing from the bodies of its rules *all* negative literals. Clearly, the program $(P_M)^h$ coincides with the *Gelfond-Lifschitz* reduct of P with respect to M (throughout the paper, we write P_M^h for $(P_M)^h$, to simplify notation). The *Gelfond-Lifschitz* operator on the algebra of all subsets of At, GL (following our convention, we omit the reference to P from the notation), is defined by

$$GL(M) = LM(P_M^h),$$

where LM(Q) stands for a least model of a Horn program Q.

We now present characterizations of the well-founded semantics. We phrase them in the language of operators and their fixpoints. All operators considered here are defined on the algebra of subsets of At(P). We denote a least fixpoint (if it exists) of an operator O by lfp(O).

It is well known that GL is antimonotone. Consequently, $GL^2 = GL \circ GL$ is monotone and has a least fixpoint. The set of atoms that are true with respect to the well-founded semantics of a program P, denoted by T_{wfs} , is precisely the least fixpoint of the operator GL^2 , that is, $T_{wfs} = lfp(GL^2)$ (Van Gelder, 1989; Fitting, 2001). The set of atoms that are false with respect to the well-founded semantics of a program P, denoted by F_{wfs} , is given by $\overline{GL(T_{wfs})}$ (throughout the paper, \overline{X} denotes the complement of a set X with respect to At(P)).

One can define a dual operator to GL^2 by

$$A(M) = \overline{GL(GL(\overline{M}))}.$$

It is easy to see that A is monotone and that its least fixpoint is F_{wfs} . Thus, $F_{wfs} = lfp(A)$ and $T_{wfs} = GL(\overline{F_{wfs}})$.

We close this section by discussing ways to compute GL(M) for a given finite propositional logic program P and a set of atoms $M \subseteq At(P)$. A straightforward approach is to compute the Gelfond-Lifschitz reduct P_M^h and then to compute its least model. However, in this paper we will use a different approach, more appropriate for the computation of the well-founded semantics. Let P be a logic program with negation. We define $At^-(P) = \{ \mathbf{not}(a) : a \in At(P) \}$. For every set $M \subseteq At(P) \cup At^-(P)$, we define $true(M) = M \cap At(P)$. If we interpret literals of $At^-(P)$ as new atoms, then for every set $M \subseteq At(P)$, the program $P \cup \mathbf{not}(M)$ can be viewed as a Horn program. Thus, it has a least model. It is easy to see that

$$GL_P(M) = true(LM(P \cup \mathbf{not}(\overline{M}))).$$

Here, P appearing at the left-hand side of the equation stands for the original logic program, while P appearing at the right-hand side of the equation stands for the same program but interpreted as a Horn program. Thus, using the algorithm of Dowling & Gallier (1984), the Gelfond-Lifschitz reduct can be computed in time O(size(P) + |M|) = O(size(P)) (since $M \subseteq At(P)$, |M| = O(size(P))).

3 Algorithms

The departure point for our discussion of algorithms to compute the well-founded semantics is the *alternating-fixpoint* algorithm of Van Gelder (1989). Using the terminology introduced in the previous section it can be formulated as follows.

```
Algorithm 1 (Van Gelder)
F := \emptyset;
repeat
T := \underbrace{true(LM(P \cup not(F));}_{F := LM(P_T^h)}; (* \text{ or equivalently: } T := GL(\overline{F}); *)
F := \overline{LM(P_T^h)}; (* \text{ or equivalently: } \overline{GL(T)}; *)
until no change in F;
return T and F.
```

Let F' and F'' be the values of the set F just before and just after an iteration of the **repeat** loop in Algorithm 1. Clearly,

$$F'' = \overline{GL(GL(\overline{F'}))} = A(F').$$

Thus, after iteration i of the **repeat** loop, $F = A^i(\emptyset)$. Consequently, it follows from our earlier remarks that when Algorithm 1 terminates, the set F that is returned satisfies $F = F_{wfs}$. Since there is no change in F in the last iteration, when the algorithm terminates, we have $T = T_{wfs}$. That is, Algorithm 1 is correct.

We will now modify Algorithm 1. The basis for Algorithm 1 is the operator A. This operator is not *progressive*. That is, M is not necessarily a subset of A(M). We will now introduce a related progressive operator, say B, and show that it can be used to replace A. Let P be a logic program and let T and F be two subsets of At(P). By $P_{F,T}$ we denote the program obtained from P by removing

- 1. all rules whose heads are in F
- 2. all rules whose bodies contain a positive occurrence of an atom from F
- 3. all rules whose bodies contain a negated literal of the form not(a), where $a \in T$.

Clearly, $P_{F,T} \subseteq P_T$.

We define an operator B(F) as follows:

$$B(F) = \overline{LM(P_{F,T}^h)},$$

where $T = GL(\overline{F})$ and $P_{F,T}^h$ abbreviates $(P_{F,T})^h$. The following result gathers key properties of the operator B.

Theorem 3.1

Let *P* be a normal logic program. Then:

- 1. B is monotone
- 2. For every $F \subseteq At(P)$, $A(F) \subseteq B(F)$
- 3. For every $F \subseteq F_{wfs}$, $B(F) \subseteq F_{wfs}$
- 4. $lfp(B) = F_{wfs}$
- 5. For every $F \subseteq At(P)$, $B(F) = F \cup (\overline{F} \setminus LM(P_{F,T}^h))$, where $T = GL(\overline{F})$.

Proof

- (1) Assume that $F_1 \subseteq F_2$. Set $T_i = GL(\overline{F_i})$, i = 1, 2. Clearly, $\overline{F_2} \subseteq \overline{F_1}$ and, by antimonotonicity of GL, $T_1 \subseteq T_2$. By the definition of $P_{F,T}$, $P_{F_2,T_2} \subseteq P_{F_1,T_1}$. Consequently, $LM(P_{F_2,T_2}^h) \subseteq LM(P_{F_1,T_1}^h)$ and, so, $B(F_1) \subseteq B(F_2)$.
- (2) Let $T = GL(\overline{F})$. Clearly, $P_{F,T} \subseteq P_T$. Thus, $A(F) = \overline{LM(P_T^h)} \subseteq \overline{LM(P_{F,T}^h)} = B(F)$.
- (3) We have, $LM(P_{T_{w/s}}^h) = \overline{F_{w/s}}$. It follows that removing from $P_{T_{w/s}}^h$ rules with heads in $F_{w/s}$ and those that contain a positive occurrence of an atom from $F_{w/s}$ in their bodies does not change the least model. That is,

$$LM(P_{F_{wfs},T_{wfs}}^h) = LM(P_{T_{wfs}}^h).$$

Since, $T_{wfs} = GL(\overline{F_{wfs}})$, $B(F_{wfs}) = \overline{LM(P_{F_{wfs}}^h, T_{wfs})}$. Let $F \subseteq F_{wfs}$. Then, by (1), $B(F) \subseteq B(F_{wfs})$. Thus, we have

$$B(F) \subseteq B(F_{wfs}) = \overline{LM(P_{F_{wfs}, T_{wfs}}^h)} = \overline{LM(P_{T_{wfs}}^h)} = F_{wfs}.$$

- (4) The least fixpoint of B is given by $lfp(B) = \bigcup B^i(\emptyset)$. By (3), $lfp(B) \subseteq F_{wfs}$. On the other hand, by (1) and (2), $A^i(\emptyset) \subseteq B^i(\emptyset)$. Thus, $F_{wfs} = lfp(A) \subseteq lfp(B)$. It follows that $lfp(B) = F_{wfs}$.
- (5) Let $T = GL(\overline{F})$. Since $P_{F,T}$ has no rules with head in F, $LM(P_{F,T}^h) \subseteq \overline{F}$ and, consequently, $F \subseteq B(F)$. Thus, the assertion follows. \square

Theorem 3.1 allows us to prove the correctness of the following modification of Algorithm 1.

```
Algorithm 2
F := \emptyset;
repeat
T := true(LM(P \cup not(F));
\Delta F := \overline{F} \setminus LM(P_{F,T}^h);
F := F \cup \Delta F;
until no change in F;
```

return T and F.

By Theorem 3.1, each iteration of the **repeat** loop computes B(F) as the new value for the set F. More formally, the set F just after iteration i, satisfies $F = B^i(\emptyset)$. Thus, when the algorithm terminates, the set F that is returned is the least fixpoint of B. Consequently, by Theorem 3.1(4), Algorithm 2 is correct.

We will now modify Algorithm 2 to obtain a general template for an alternating-fixpoint algorithm to compute the well-founded semantics. The key idea is to observe that it is enough to compute a subset of ΔF in each iteration and the algorithm remains correct.

Let us assume that for some operator Δ_w defined for pairs (F,Q), where $F \subseteq At(P)$ and Q is a Horn program such that $At(Q) \subseteq \overline{F}$ (the complement is, as always, evaluated with respect to At(P)), we have:

(W1)
$$\Delta_w(F,Q) \subseteq \overline{F} \setminus LM(Q)$$
 (W2) $\Delta_w(F,Q) = \emptyset$ if and only if $\overline{F} \setminus LM(Q) = \emptyset$.

Let $F \subseteq At(P)$. By the definition of $P_{F,T}$, $At(P_{F,T}^h) \subseteq \overline{F}$. Thus, we define $B_w(F) = F \cup \Delta_w(F, P_{F,T}^h)$, where $T = true(LM(P \cup \mathbf{not}(F)))$. It is clear that for every $F \subseteq At(P)$, $F \subseteq B_w(F) \subseteq B(F)$, the latter inclusion follows from Theorem 3.1(5) and (W1). Consequently, for every i,

$$B_w^i(\emptyset) \subseteq B^i(\emptyset).$$

It follows that $B_w^i(\emptyset) \subseteq lfp(B) = F_{wfs}$. It also follows that there is the first i such that $B_w^i(\emptyset) = B_w^{i+1}(\emptyset)$. Let us denote this set $B_w^i(\emptyset)$ by F_0 . Then $F_0 \subseteq F_{wfs}$. In the same time, by condition (W2), $B(F_0) = F_0$. Since F_{wfs} is the least fixpoint of B, $F_{wfs} \subseteq F_0$. It follows that a modification of Algorithm 2 in which line

$$\Delta F := \overline{F} \setminus LM(P_{F,T}^h);$$

is replaced by

$$\Delta F := \Delta_w(F, P_{FT}^h);$$

correctly computes the well-founded semantics of a program P. Thus, we obtain the following algorithm for computing the well-founded semantics.

```
Algorithm 3
F := \emptyset;
repeat
T := true(LM(P \cup not(F));
\Delta F := \Delta_w(F, P_{F,T}^h);
F := F \cup \Delta F;
until no change in F;
return T and F.
```

We will now refine Algorithm 3. Specifically, we will show that the sets T and F can be computed incrementally.

Let R be a Horn program. We define the *residual* program of R, res(R), to be the Horn program obtained from R by removing all rules of R with the head in LM(R) and by removing from the bodies of the remaining rules those elements that are in LM(R). We have the following technical result.

Lemma 3.2

Let *R* be a Horn program and let *M* be a set of atoms such that $M \cap head(R) = \emptyset$. Then $LM(R \cup M) = LM(R) \cup LM(res(R) \cup M)$.

Lemma 3.2 implies that (we treat here negated literals as new atoms and P as Horn program over the extended alphabet)

$$LM(P \cup \mathsf{not}(F \cup \Delta F)) = LM(P \cup \mathsf{not}(F)) \cup LM(res(P \cup \mathsf{not}(F)) \cup \mathsf{not}(\Delta F)).$$

Thus, if the set F is expanded by new elements from ΔF , then the new set T can be computed by increasing the old set T by $\Delta T = true(LM(res(P \cup \mathbf{not}(F)) \cup \mathbf{not}(\Delta F)))$. Important thing to note is that the increment ΔT can be computed on the basis of the residual program and the increment ΔF . Similarly, we have

$$P_{F \cup \Delta F, T \cup \Delta T} = (P_{F,T})_{\Delta F, \Delta T}.$$

Thus, computing $P_{F,T}$ can also be done incrementally on the basis of the program considered in the previous iteration by taking into account most recently computed increments ΔF and ΔT .

This discussion implies that Algorithm 3 can be equivalently restated as follows:

```
Algorithm 3
      T := F := \Delta T := \Delta F := \emptyset;
1
2
      R := P; (*R will be treated as a Horn program *)
3
      Q := P;
4
      repeat
5
         \Delta T := true(LM(R \cup \mathbf{not}(\Delta F));
6
         R := res(R \cup not(\Delta F));
7
         T := T \cup \Delta T;
8
         Q := Q_{\Delta F, \Delta T};
9
         \Delta F := \Delta_w(F, Q^h);
10
         F := F \cup \Delta F;
      until no change in F;
11
     return T and F.
```

We will now estimate the running time of Algorithm 3. Clearly line 1 requires a constant time. Setting up appropriate data structures for programs R and Q (lines 2 and 3) takes O(size(P)) steps. In each iteration, ΔT is computed and the current program R is replaced by the program $res(R \cup not(\Delta F))$ (lines 5 and 6). By modifying the algorithm from Dowling & Gallier (1984), and assuming that R is already stored in the memory (it is avaliable either as the result of the initialization in the case of the first iteration or as a result of the computation in the previous iteration), both tasks can be accomplished in $O(size(R^o) + |\Delta F| - size(R^n))$ steps. Here R^o denotes the old version of R and R^n denotes the new version of R. Consequently, the total time needed for lines 5 and 6 over all iterations is given by $O(size(P) + |At(P)| - size(R^t)) = O(size(P))$ (where R^t is the program R, when the algorithm terminates). The time needed for all executions of line 7 is proportional to the number of iterations and is O(|At(P)|) = O(size(P)).

Given a logic program Q and sets of atoms ΔT and ΔF , it takes $O(size(Q) - size(Q_{\Delta F,\Delta T}) + |\Delta T| + |\Delta F|)$ steps to compute the program $Q_{\Delta F,\Delta T}$ in line 8. We assume here that Q is already in the memory as a result of the initialization in the case of the first iteration, or as the result of the computation in the previous iteration, otherwise. It follows that the total time over all iterations needed to execute line 8 is O(size(P) + |At(P)|) = O(size(P)).

Thus, we obtain that the running time of Algorithm 3 is given by O(size(P) + m), where m is the total time needed to compute $\Delta_w(F, Q^h)$ over all iterations of the algorithm.

In the standard (Van Gelder's) implementation of Algorithm 3, we compute the whole set $\overline{F} \setminus LM(Q^h)$ as $\Delta_w(F,Q^h)$. In addition, computation is performed in a bottom-up fashion. That is, we first compute the least model of Q^h and then its complement with respect to \overline{F} . Such approach requires $O(size(Q^h)) = O(size(P))$

steps per iteration to execute line 9 and leads to $O(|At(P)| \times size(P))$ running-time estimate for the alternating-fixpoint algorithm.

4 Procedure Δ_w

In this section we will focus on the class of programs, \mathcal{LP}_1 , that is, programs whose rules have no more than one positive atom in their bodies. We assume that we have a procedure *false* that, given a Horn program $Q \in \mathcal{LP}_1$, returns a subset of the set $At(Q) \setminus LM(Q)$. We also assume that *false* returns the empty set *if and only if* At(Q) = LM(Q). For every pair (F,Q), where $F \subseteq At(P)$ and Q is a Horn program such that $At(Q) \subseteq \overline{F}$, we define

$$\Delta_w(F,Q) = false(Q).$$

It is easy to see that this operator $\Delta_w(F,Q)$ satisfies conditions (W1) and (W2). Consequently, it can be used in Algorithm 3. Clearly, the procedure Δ_w and its computational properties are determined by the procedure *false*. In the remainder of the paper, we will describe a particular implementation of the procedure *false* and estimate its running time. We will use this estimate to obtain a bound on the running time of the resulting version of Algorithm 3.

A straightforward way to compute the least model of Q and so, to find $At(Q) \setminus LM(Q)$, is 'bottom-up'. That is, we start with atoms which are heads of rules with the empty bodies and use the rules of Q to compute all atoms in LM(Q) by iterating the van Emden-Kowalski operator. An efficient implementation of the process is provided by the Dowling-Gallier algorithm (Dowling & Gallier, 1984).

The approach we follow here in the procedure *false* is "top-down" and gives us, in general, only a part of the set $At(Q) \setminus LM(Q)$. More precisely, for an atom a we proceed 'backwards' attempting to construct a proof or to demonstrate that no proof exists. In the process, we either go back to an atom that is the head of a rule with empty body or we show that no proof exists. In the former case, $a \in LM(Q)$. In the latter one, none of the atoms considered while searching for a proof of a are in LM(Q) (because $Q \in \mathcal{LP}_1$ and each rule has at most one antecedent). The problem is that we may find an atom a that does not have a proof only *after* we look at all other atoms first. Thus, in the worst case, finding one new false atom may require time that is proportional to the size of Q.

To improve the time performance, we look for proofs simultaneously for all atoms and grow the proofs 'backwards' in a carefully controlled way. Namely, we never let one search to get too much ahead of the other searches. This controlled way of looking for proofs is the key idea of our approach and leads to a better performance. We will now provide an informal description of the procedure *false* followed later by a formal specification and an example.

In the procedure, we make use of a *new* atom, say s, different from all atoms occurring in Q. Further, we denote by head(r) the atom in the head of a rule $r \in Q$ and by tail(r) the atom which is either the unique positive atom in the body of r, if such an atom exists, or s otherwise. We call an atom $a \in At(Q)$ accessible if there are rules r_1, \ldots, r_k in Q such that $tail(r_{i+1}) = head(r_i)$, for $i = 1, \ldots, k-1$, $tail(r_1) = s$

and $head(r_k) = a$. Clearly, the least model LM(Q) of Q is precisely the set of all accessible atoms.

In each step of the algorithm, the set of atoms from At(Q) is partitioned into potentially false sets or pf-sets, for short. We say that a set $v \subseteq At(Q)$ is a pf-set if for each pair of distinct atoms $a, b \in v$ there are rules r_1, \ldots, r_k in Q such that $tail(r_{i+1}) = head(r_i) \in v$, for $i = 1, \ldots, k-1$, $tail(r_1) = b$ and $head(r_k) = a$. It is clear that if v is a pf-set then either all its elements are accessible (belong to the least model of Q) or none of them does (they are all false). Clearly, singleton sets consisting of individual atoms in At(Q) are pf-sets. In the algorithm, with each pf-set we maintain its cardinality.

Current information about the state of all top-down searches and about the dependencies among atoms, that were discovered so far, is maintained in a directed graph \mathscr{G} . The vertex set of this graph, say \mathscr{S} , consists of $\{s\}$ and of a family of pf-sets forming a partition of the set At(Q). The edges of \mathscr{G} are specified by a partial function $pred: \mathscr{S} \to \mathscr{S}$. We write $pred(v) = \mathbf{undefined}$ if pred is undefined for v. Thus, the set of edges of \mathscr{G} is given by $\{(pred(v), v): pred(v) \neq \mathbf{undefined}\}$. Since pred is a partial function, it is easy to see that the connected components of the graph \mathscr{G} are unicyclic graphs or trees rooted in those vertices v for which pred(v) is undefined. Throughout the algorithm we always have $pred(\{s\}) = \mathbf{undefined}$. Thus, the connected component of \mathscr{G} containing $\{s\}$ is always a tree and $\{s\}$ is its root.

If w and v are two different pf-sets, the existence of the edge (w,v) in $\mathscr G$ means that we have already discovered a rule in the original program whose head is in v and whose tail is in w. Thus, if vertices in w are accessible, then so are the vertices in v. A pf-set that is the root of a tree forming a component of $\mathscr G$ is called an *active* pf-set. If v is an active pf-set then no rule v with v with v is an active pf-set detected so far. Thus, v is a candidate for a set of atoms which does not intersect the least model of v. Let us note that even though v is a root of a tree in v it is never active as it is not a pf-set in the first place.

We let active pf-sets grow by gluing them with other pf-sets. However, we allow to grow only these active pf-sets whose cardinalities are the least. In each iteration of the algorithm the value of the variable size is a lower bound for the cardinalities of active pf-sets. To grow an active pf-set v, we look for rules with heads in v and with tails in pf-sets other than v (not necessarily active) or in $\{s\}$. The dependencies between pf-sets discovered in this way are represented as new directed edges in \mathcal{G} . Pf-sets that appear in the same cycle are glued together (in the procedure cycle). Since $\{s\}$ is not an active pf-set, it never becomes an element of a cycle in \mathcal{G} .

If, when attempting to grow a pf-set v, we discover a rule with head in v and with the tail in a vertex of the tree of \mathcal{G} rooted in $\{s\}$, then v is from now on ignored (all its vertices belong to the least model of Q). Indeed, v gets connected to a tree of \mathcal{G} rooted in $\{s\}$. Consequently, it cannot become a member of a cycle in \mathcal{G} in the future and is never again considered by the procedure cycle.

The main loop (lines 6-23) of the algorithm *false* below starts by incrementing *size* followed by a call to the procedure $cycle(\mathcal{S}, pred, size, L)$. This procedure scans the graph \mathcal{G} and identifies all its cycles. It then modifies \mathcal{G} by considering each cycle and by gluing its pf-sets into a single pf-set. To this end, it modifies the vertex set \mathcal{S}

of \mathscr{G} and the function *pred* defining the edges of \mathscr{G} . Each such new pf-set becomes the root of its tree in \mathscr{G} and so, it becomes active. The procedure *cycle* computes the cardinality of each new active pf-set. Finally, it creates a list L so that it consists of active pf-sets of cardinality *size*. If no such set is found (L is empty), we move on to the next iteration of the main loop and increment *size* by 1. We give a more detailed description of the procedure *cycle* later in the paper when we analyse the time complexity of our method.

For each active pf-set $v \in L$ we consider the tail of each rule with head in v (lines 9-22). If there is a rule r with $head(r) \in v$ and $tail(r) \notin v$ then it is detected (line 15). The value pred(v) is set to this element in $\mathscr S$ that contains tail(r) (it may be that this set is $\{s\}$). We also set the variable *success* to **true** (line 16). The pf-set v stops to be active. We move on to the next active pf-set on L.

If such a rule r does not exist then $success = \mathbf{false}$ and v is a set of cardinality size consisting of atoms which are not in the least model of Q. This set is returned by the procedure false (line 21). Hence, for an active pf-set considered in the loop 6-23, either we find a pf-set $pred(v) \in \mathcal{S} \setminus \{v\}$ (and we have to consider the next pf-set on L) or v is returned as a set of atoms which are not in the least model of Q (and the procedure false terminates). Thus, the procedure false is completed if either a nonempty set v of atoms which are not in the least model of Q is found or, if after some passes of the loop 6-23, the graph \mathcal{G} has no active pf-sets. In the latter case \mathcal{G} is a tree with the root in $\{s\}$. Thus, At(Q) = LM(Q) and $v = \emptyset$ is returned (line 24).

In the procedure *false*, as formally described below, an input program Q is represented by lists IN(a), $a \in At(Q)$, of all atoms b such that b is the body of some rule with the head a. If there is a rule with the head a and empty body, we insert s into the list IN(a).

We also use an operation next on lists and elements. Let l be a list and w be an element, either belonging to l or having a special value **undefined**. Then

```
next(w, l) = \begin{cases} \text{ the next element after } w \text{ in } l & \text{if } w \in l \\ \text{ the first element in } l & \text{if } w \text{ is } \mathbf{undefined}. \end{cases}
```

The value undefined should not be mixed with nil which indicates the end of a list.

Finally, we use a procedure $findset(w, \mathcal{S})$ which, for an atom w and a collection \mathcal{S} of disjoint sets, one of which contains w, finds the name of the set in \mathcal{S} containing w (it follows from our assumptions that such a set is unique). Elements of \mathcal{S} are maintained as linked lists. Each element on such a list has a pointer to the head of the list. The head serves as the identifier for the list. When the procedure $findset(w, \mathcal{S})$ is called, it returns the head of the list to which w belongs.

```
1 procedure false(Q);

2 \mathscr{S} := \{\{x\} : x \in At(Q)\} \cup \{\{s\}\};

3 for v \in \mathscr{S} do pred(v) := undefined;

4 for x \in At(Q) do \{w(x) := undefined; cardinality(\{x\}) := 1\};

5 size := 0;

6 while size < |At(Q)| do

7 \{size := size + 1;
```

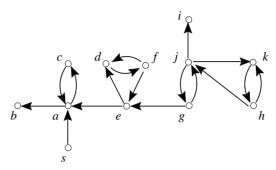


Fig. 1. Graph G^Q representing program Q.

```
8
           cycle(\mathcal{S}, pred, size, L);
9
           for all v \in L do
10
              \{success := false;
              u := next(\mathbf{undefined}, v); (* find first element on list v *)
11
12
              while u \neq \text{nil} and not success do
13
                w(u) := next(w(u), IN(u));
14
                while w(u) \neq \text{nil} and not success do
15
                   {if findset(w(u), \mathcal{S}) \neq v
                      then \{success := true; pred(v) := findset(w(u), \mathcal{S})\}\
16
17
                      else w(u) := next(w(u), IN(u))
18
                end while (14);
19
                if not success then u := next(u, v)
20
              end while (12);
21
             if not success then return v (* the procedure terminates *)
22
           end for (9)}
23
        end while (6)};
        return v = \emptyset
24
25
     end false;
```

We will now illustrate the operation of the algorithm. Let us consider the following Horn logic program Q:

This program is represented as a graph, G^Q , in Figure 1. The vertices of this graph correspond to the atoms of the program. In addition, G^Q has an auxiliary vertex $s \notin At(Q)$. An edge (x, y), where $x, y \in At(Q)$, represents the clause $y \leftarrow x$ from Q. An edge (s, y), where $y \in At(Q)$, represents the clause $y \leftarrow x$. When illustrating the algorithm, we assume that atoms from At(Q) (atoms a, \ldots, k in our example) appear on the lists IN(x), $x \in At(Q)$, in the alphabetical order. We also assume that whenever s belongs to a list IN(x), it appears as the first atom on the list.

In the algorithm false, the current state of knowledge about the possibility of

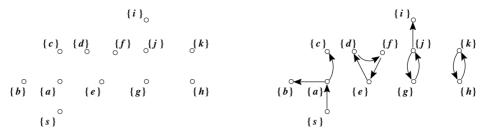


Fig. 2. Graph $\mathscr G$ initially (left side) and after the first iteration of the loop 6-23 (on the right).

proving an atom from Q is represented by the graph \mathcal{G} . Initially, \mathcal{G} consists of isolated vertices. Indeed, line 3 of the algorithm sets pred(x) to undefined, for every vertex x of \mathcal{G} (see Figure 2 (left)). All of the vertices of \mathcal{G} , except for $\{s\}$ are active pf-sets. The procedure cycle (line 8), called with size=1, puts all of them on the list L.

The algorithm considers next (line 9) all elements on the list L, that is, all vertices of G that are active pf-sets and have cardinality equal to size. During the first iteration of the loop 6-23, L consists of all vertices of \mathcal{G} , except for $\{s\}$ (that is, singleton sets $\{x\}$, where $x \in At(Q) = V(G^Q) \setminus \{s\}$). For each vertex v of \mathscr{G} on L, the algorithm looks for a back rule for v, that is, a rule in Q with the head in v and the tail in a pf-set other than v or in $\{s\}$. In our graphical representation of Q by means of the graph G^Q , a back rule for v corresponds to an edge (referred to as a back edge) in G^Q with the head in v and the tail in a vertex of $\mathscr G$ other than v (possibly in $\{s\}$). To find a back rule (edge) for v, all atoms u of Q (equivalently, all vertices u of G^Q) that belong to v are considered (the loop 12-20). For each such atom u, the algorithm searches for the first atom on the list IN(u) that does not belong to v. Let us recall that IN(u) is the list of atoms that are the tails of rules with the head u or, in the terms of the graph G^{Q} , that are the tails of edges with the head u. If such an atom is found, together with u it determines a back rule (edge) r for v. The algorithm sets pred(v) to be equal to the pf-set containing the tail of r (line 16). That is, an edge from pred(v) to v is added to \mathcal{G} . The algorithm moves then on to the next element of the list L.

In our example, in the first iteration of the loop 6-23, a back rule is found for every element on L, that is, for every vertex of \mathcal{G} other than $\{s\}$. For instance, for the vertex $\{d\}$, the algorithm considers atoms on the list IN(d) = (e, f) (let us recall that atoms on lists IN(x) are arranged alphabetically with the exception of the special atom s which, if present on a list, is always its first element). The first atom on the list, e does not belong to $\{d\}$. Thus, it defines, together with d a back rule for $\{d\}$, $d \leftarrow e$. The resulting graph \mathcal{G} is shown in Figure 2 on the right.

Let us note that when scanning the list IN(d) in subsequent iterations the algorithm resumes the scan with the first atom that has not been looked at yet (cf. the definition of the operation next). Thus, the next time d is considered as an element of an active pf-set for which a back rule is searched for, the scan of IN(d) will start with f. The same holds true for all lists IN(x), $x \in At(Q)$. Consequently, each atom on each of these lists is considered just once. Such an approach still guarantees that

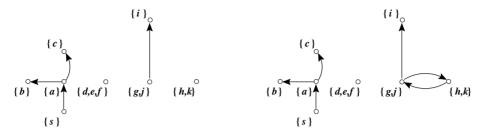


Fig. 3. Graph \mathscr{G} after the execution of the procedure *cycle* in the second iteration of the loop 6-23 (left) and after the second iteration of the loop 6-23 (right).

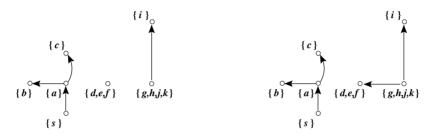


Fig. 4. Graph \mathscr{G} after the execution of the procedure *cycle* in the third iteration of the loop 6-23 (left) and after the third iteration of the loop 6-23 (right).

finding back rules works correctly (that is, that they are found by the algorithm whenever they exist). Indeed, when an atom on a list IN(x) is considered, it either defines a back rule with the head x (and thus cannot define any new back rule with the head x in the future) or it is in the same active pf-set as x (and thus it neither defines a back rule now nor it will define it in the future, as it will remain in the same pf-set as x till the algorithm terminates).

The second iteration of the loop 6-23 starts with the procedure *cycle* contracting each cycle in the graph \mathcal{G} to a single vertex. The resulting graph is shown in Figure 3 on the left. The procedure *cycle* then creates a new list L. It consists of all active pf-sets of cardinality 2. In our case, L contains $\{g, j\}$ and $\{h, k\}$ ($\{d, e, f\}$ is also active but has cardinality 3).

Continuing with the second iteration, the algorithm next considers each vertex on L (the loop 9-22) and looks for back rules. In this iteration, a back rule is found for each of the nodes on L and the modified graph \mathcal{G} is given in Figure 3 on the right.

In the third iteration, the procedure *cycle* contracts the only cycle in \mathscr{G} to a single active pf-set of cardinality 4 (Figure 4, left side). It also creates a new list L. This time it consists of active pf-sets of cardinality 3. There is just one such set $-\{d,e,f\}$. Subsequently, the algorithm *false* looks for a back rule for $\{d,e,f\}$. It starts by considering edges ending in d (line 11; we assume that v is represented by the list (d,e,f)). It scans the list IN(d) starting at the first atom that has not been inspected so far, i.e f. However, since f belongs to the same pf-set as d, f does not specify a back rule. Since there are no more atoms on the list IN(d), we move on to the next iteration of the loop 12-20 and consider atom e. We have IN(e) = (f,g). Since f was already considered (and yielded a back rule for $\{e\}$) in the first iteration, we consider g. Since $g \notin \{d,e,f\}$, it defines a back rule for $\{d,e,f\}$, $e \leftarrow g$.

The resulting graph \mathcal{G} is shown in Figure 4 (on the right). It has no cycles. So, the only thing done by the procedure *cycle* in the iteration 4 is that it puts on L active pf-sets of cardinality 4. There is just one such set in \mathcal{G} , $\{g,h,j,k\}$. The algorithm *false* looks for a back edge for $\{g,h,j,k\}$ and does not find any. The variable *success* remains **false**. The algorithm returns $\{g,h,j,k\}$ and terminates (line 21). Let us note that this set is a proper subset of the set $At(Q) \setminus LM(Q)$.

The following theorem formally establishes two key properties of the procedure false.

Theorem 4.1

- 1. The procedure *false* returns a set v such that $v \subseteq At(Q) \setminus LM(Q)$.
- 2. false returns the empty set if and only if $At(Q) \setminus LM(Q) = \emptyset$.

Proof

(1) The statement is trivially true if *false* returns the empty set. Thus assume that the returned set $v \neq \emptyset$. It means that the value of the variable *success* is **false** after all passes of the loop 12-20 for some active pf-set v in the list L. Thus every rule in Q with the head in v has been considered.

Suppose there is a rule r in Q with $head(r) = u \in v$ and $tail(r) = b \notin v$. This rule was considered by the procedure false when u = head(r) was a member of some active pf-set, say y. Since larger pf-sets are obtained by gluing smaller ones, $y \subseteq v$. While r was being considered, the value of w(u) in the loop 14-18 was b and the value of v was v. Consequently, $findset(b, \mathcal{S}) \neq v$ in line 15 because $v \subseteq v$ and $v \notin v$ so $v \notin v$. Hence the value of $v \in v$ says set to $v \in v$ stopped to be active. Recall that $v \in v$ is active when the procedure stops. Hence $v \in v$ had to be glued with other pf-sets to obtain $v \in v$. This is, however, impossible because if $v \in v$ were glued with some other pf-sets to form a larger pf-set $v \in v$. We have got a contradiction with $v \in v \in v$.

Hence, there are no rules r in Q with $head(r) \in v$ and $tail(r) \notin v$. Thus no atom in v is accessible so $v \subseteq At(Q) \setminus LM(Q)$.

(2) Suppose false returns the empty set and consider the last pass of the loop 6-23, for size = |At(Q)|. If the list L is empty then no vertex of $\mathscr G$ is an active pf-set. Hence, $\mathscr G$ is a tree with the root $\{s\}$. Thus all atoms in At(Q) are accessible and consequently LM(Q) = At(Q).

If the list L is nonempty then it contains one pf-set v = At(Q). The empty set is returned by the procedure *false* so the value of the variable *success* in line 16 is **true** for v = At(Q). It means that for some rule r in Q with head(r) = u, $w(u) = tail(r) \notin v = At(Q)$ so w(u) = s. Hence, u is accessible and, consequently, all atoms in At(Q) are accessible. That is, we have $At(Q) \setminus LM(Q) = \emptyset$.

The converse of the implication proved above follows immediately from the first part of the theorem. \Box

We shall now consider the procedure *cycle* a little bit more carefully. The procedure can be informally written in the following form:

procedure $cycle(\mathcal{S}, pred, size, L)$

- 1. Initialize L to empty.
- 2. Find all cycles $C_1, C_2, ..., C_p$ in the graph \mathscr{G} . Put $\mathscr{C} = \{C_1, C_2, ..., C_p\}$.
- 3. For every cycle $C = \{v_1, \dots, v_q\}, C \in \mathcal{C}$, do (i)-(iv).
 - (i) set $v_C := v_1 \cup \ldots \cup v_q$;
 - (ii) compute $cardinality(v_C)$ (sum up the cardinalities of all vertices in C);
 - (iii) update the set \mathcal{S} : set $\mathcal{S} := (\mathcal{S} \{v_1, \dots, v_q\}) \cup \{v_C\}$; (* v_C becomes an active pf-set *)
 - (iv) update the function *pred*: for every i = 1, ..., q, if $pred(z) = v_i$ (for some $z \in \mathcal{S}$) then $pred(z) := v_C$;
- 4. For every vertex of \mathcal{G} that is an active pf-set, if cardinality(v) = size, insert v into the list L.

Since \mathscr{G} is a directed graph whose connected components are either unicyclic graphs or trees, step 2 of the procedure cycle can be implemented in $O(|\mathscr{S}|)$ time. Since pf-sets are represented as linked lists, with each node on the list pointing to the head of the list, step (i) can be implemented to take $O(|v_C|)$ steps. The time needed for step (ii) is, clearly, O(|C|). Each execution of step (iii) takes also O(|C|). Finally, the running time of each execution of step (iv) is $O(m_C)$, where m_C is the size of the connected component of the graph \mathscr{G} containing C. Thus, an iteration of the loop 3 for a cycle $C \in \mathscr{C}$ takes $O(|C| + m_C + |v_C|)$. Clearly, $|C| \leq m_C$. Moreover, $\sum_{C \in \mathscr{C}} m_C \leq |\mathscr{S}| - 1 \leq |At(Q)|$ and $\sum_{C \in \mathscr{C}} |v_C| \leq |At(Q)|$ (they are all disjoint subsets of At(Q)). Thus, the total time needed for the loop 3 is O(|At(Q)|). It is easy to see that the time needed for the loop 4 is also O(|At(Q)|). Consequently, the running time of the procedure cycle is O(|At(Q)|).

We are now in a position to estimate the running time of the procedure false.

Lemma 4.2

If the procedure false(Q) returns a nonempty set v, then the running time of false is $O(|v| \times |At(Q)|)$. If false(Q) returns the empty set then its running time is $O(|At(Q)|^2)$.

Proof

Let |At(Q)| = n and |v| = k. As we have already observed the procedure *cycle* runs in time O(n). It is not hard to see that, since we represent all sets occurring in the procedure *false* as linked lists, with each node on a list pointing to the head of the list, the operations: *findset* and *next* require a constant time.

First assume that the output v of the procedure *false* is nonempty. Let us estimate the number of passes of the **while** and **for** loops in the procedure. Clearly, the loop 6-23 is executed k times. Hence the total running time of all calls of the procedure cycle is O(kn). The number of passes of the loop 9-22 is not larger than $|L_1| + |L_2| + \ldots + |L_k|$, where L_i denotes the list L in an iteration i of the loop. Since L_i is a list of disjoint pf-sets of cardinality i, $|L_i| \le n$, for each $i = 1, 2, \ldots, k$. Hence the number of passes of the loop 9-22 can be very roughly estimated by kn. The loop 12-20 is executed at most

$$\sum_{i=1}^{k} \sum_{v \in L_i} |v| \leqslant kn$$

times. This inequality follows from the fact that the sets v in the lists L_i are disjoint subsets of atoms so $\sum_{v \in L_i} |v| \le n$. The estimation of the number of passes of the loop 14-18 is a little bit more complicated. First notice that in each execution of the loop we check a rule of the program Q and rules are checked only one time. The rules r checked in the loop have either both the head and the tail in some pf-set $v \in \mathscr{S}$ or $head(r) \in v$ and tail(r) is in some other pf-set $u \in \mathscr{S}$. In the latter case pred(v) is defined in line 16. The number of executions of line 16 is not larger than the number of passes of the loop 9-22 so it is bounded by kn. When the procedure returns the output, the pf-sets have cardinalities not larger than k. Hence the number of rules with both the head and the tail in the same pf-set that has been checked before the procedure stops is not larger than

$$\sum_{u \in \mathcal{S}} |u|(|u|-1) \leqslant (k-1) \sum_{u \in \mathcal{S}} |u| \leqslant (k-1)n.$$

Thus the number of passes of the loop 14-18 in the whole procedure *false* is less than 2kn. It follows that if the output v of *false* is nonempty then the running time of *false* is $O(|v| \times |At(Q)|)$.

Now consider the case when the procedure *false* returns the empty set. Clearly the number of passes of the loop 6-23 is n so it takes $O(n^2)$ time for all executions of the procedure *cycle*. Since the rules are checked in the loop 14-18 only one time, the number of passes of this loop is not larger than the number m of rules in Q. Obviously $m \le n^2$ so the running time of *false* in this case is $O(|At(Q)|^2)$.

By Lemma 4.2 and considerations in Section 3 we get an estimation of the running time of Algorithm 3.

Theorem 4.3

If P is a program whose rules have at most one positive atom in the body then Algorithm 3 can be implemented so that its running time is $O(|At(P)|^2 + size(P))$. \square

5 Conclusions

The method for computing the well-founded semantics described in this paper is a refinement of the basic alternating-fixpoint algorithm. The key idea is to use a top-down search when identifying atoms that are false. Our method is designed to work with programs whose rules have at most one positive atom in their bodies (class \mathcal{LP}_1). Its running time is $O(|At(P)|^2 + size(P))$ (where P is an input program). Thus, our algorithm is an improvement over other known methods to compute the well-founded semantics for programs in the class \mathcal{LP}_1 . Our algorithm runs in linear time for the class of programs $P \in \mathcal{LP}_1$ for which $size(P) \ge |At(P)|^2$. However, it is not a linear-time algorithm in general. It is an open question whether a linear-time algorithm for computing the well-founded semantics for programs in the class \mathcal{LP}_1 exists.

Our results extend to the class \mathcal{LP}_1^+ . However, the extension is straightforward and the class \mathcal{LP}_1^+ is still rather narrow. Moreover, it is not specified syntactically (it is described by means of the Kripke-Kleene semantics). The question arises whether

our top-down approach to positive-loop detection can be generalized to any class of programs significantly extending the class \mathcal{LP}_1 and possessing a simple syntactic description.

Finally, let us note that the general problem of computing the well-founded semantics still remains a challenge. No significant improvement over the alternating-fixpoint algorithm of Van Gelder has been obtained for the class of arbitrary finite propositional logic programs.

Acknowledgments

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