# Solving Horn Clauses on Inductive Data Types Without Induction

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submitted 1 January 2003; revised 1 January 2003; accepted 1 January 2003

## Abstract

We address the problem of verifying the satisfiability of Constrained Horn Clauses (CHCs) based on theories of inductively defined data structures, such as lists and trees. We propose a transformation technique whose objective is the removal of these data structures from CHCs, hence reducing their satisfiability to a satisfiability problem for CHCs on integers and booleans. We propose a transformation algorithm and identify a class of clauses where it always succeeds. We also consider an extension of that algorithm, which combines clause transformation with reasoning on integer constraints. Via an experimental evaluation we show that our technique greatly improves the effectiveness of applying the Z3 solver to CHCs. We also show that our verification technique based on CHC transformation followed by CHC solving, is competitive with respect to CHC solvers extended with induction.

This paper is under consideration for acceptance in TPLP.

*KEYWORDS*: Program verification, constrained Horn clauses, constraint logic programming, inductively defined data types, program transformation.

#### 1 Introduction

Constraint logic programs have become a well-established formalism for solving program verification problems (Albert et al. 2007, Bjørner et al. 2015, De Angelis et al. 2014, Jaffar et al. 2012, Méndez-Lojo et al. 2008, Peralta et al. 1998). In the verification field, constraint logic programs are often called constrained Horn clauses (CHCs), and here we will adopt this terminology. The verification method based on CHCs consists in reducing a program verification problem to the satisfiability of a set of CHCs. Since CHC satisfiability, also called CHC solving, is in general an undecidable problem, some heuristics have been proposed in the literature, such as *Counterexample Guided Abstraction Refinement* (CEGAR) (Clarke et al. 2000), *Craig interpolation* (McMillan 2003), and *Property Directed Reachability* (PDR) (Bradley 2011, Hoder and Bjørner 2012). Some

tools, called CHC solvers, for verifying the satisfiability of CHCs are available. We recall Eldarica (Hojjat et al. 2012), HSF (Grebenshchikov et al. 2012), RAHFT (Kafle et al. 2016), VeriMAP (De Angelis et al. 2014), and Z3 (de Moura and Bjørner 2008). CHC solvers make use of a combination of the above heuristics and have been shown to be very effective for CHCs with several underlying constraint theories, such as the theory of Linear Integer Arithmetic (LIA) and the theory of Boolean constraints (Bool).

Solving techniques have also been developed for CHCs manipulating inductively defined data structures such as lists and trees (see, for instance, https://rise4fun.com/Z3/ tutorial/guide for the Z3 solver). However, CHC solvers acting on those data structures are usually less effective than CHC solvers for clauses with constraints in *LIA* or *Bool*. This is mainly due to the fact that methods for satisfiability used by CHC solvers are based on variants of resolution, augmented with ad hoc algorithms for the underlying constraint theory, and no induction principles for the data structures are used.

To overcome this difficulty, recent work has proposed the extension of CHC solving by adding a principle of induction on predicate derivations (Unno et al. 2017). This work is on the same line of other proposals which extend techniques for Satisfiability Modulo Theory (SMT) with inductive reasoning (Reynolds and Kuncak 2015, Suter et al. 2011) by incorporating methods derived from the field of automated theorem proving.

In this paper we propose an alternative method to solve CHCs on inductively defined data structures. It is based on the application of suitable transformations of CHCs that have the objective of removing those data structures while preserving satisfiability of clauses. Our transformation method makes use of the fold/unfold transformation rules (Etalle and Gabbrielli 1996, Tamaki and Sato 1984), and it is based on some techniques proposed in the past for improving the efficiency of functional and logic programs, such as *deforestation* (Wadler 1990), *unnecessary variable elimination* (Proietti and Pettorossi 1995), and *conjunctive partial deduction* (De Schreye et al. 1999). However, the focus of the method presented in this paper is on the improvement of effectiveness of CHC solvers, rather than the improvement of efficiency of programs.

In our method we separate the concern of reasoning on inductively defined data structures from the concern of proving clause satisfiability. This separation of concerns eases the task of the CHC solvers in many cases. For instance, when the constraints are on trees of integers, our transformation method, if successful, allows us to derive an equisatisfiable set of clauses with constraints on integers only.

The main contributions of our work are the following. (1) We propose a transformation algorithm, called  $\mathcal{E}$ , that removes inductively defined data structures from CHCs and derives new, equisatisfiable sets of clauses whose constraints are from the *LIA* theory and the *Bool* theory (see Section 4). (2) We identify a class of CHCs where Algorithm  $\mathcal{E}$  is guaranteed to terminate, and all inductively defined data structures can be removed (see Section 5). (3) By using reasoning techniques on *LIA* constraints, we derive an extension of our transformation algorithm  $\mathcal{E}$ , called Algorithm  $\mathcal{EC}$  (see Section 6). (4) We report on an experimental evaluation of a tool that implements Algorithm  $\mathcal{EC}$ . We consider CHCs that encode properties of OCaml functional programs, and we show that our tool is competitive with the RCAML tool that extends the Z3 solver by adding induction rules for reasoning on inductive data structures (Unno et al. 2017) (see Section 7).

# 2 A Tree Processing Example

As an introductory example to illustrate our technique for proving program properties without using induction on data structures, let us consider the following *Tree\_Processing* program, which we write according to the OCaml syntax (Leroy et al. 2017).

 $\begin{array}{l} \mbox{type tree} = \mbox{Leaf} \mid \mbox{Node of } int * tree * tree ;; \\ \mbox{let } min \; x \; y = \mbox{if } x < y \; \mbox{then } x \; \mbox{else } y \; ;; \\ \mbox{let rec } min\_leaf \; t = \mbox{match } t \; \mbox{with} \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & &$ 

| Node $(x,l,r) \rightarrow if n \le 0$  then Node(x,l,r) else  $left_drop(n-1) l;;$ 

Let us also consider the following non-trivial property *Prop* to be verified for the *Tree\_Processing* program:

 $\forall n \; \forall t. \; n \ge 0 \; \Rightarrow \; \left( \left( (\min_{l \in f_{l} drop \; n \; t}) + n \right) \ge \left( \min_{l \in f_{l} t} t \right) \right).$ 

Now we translate the *Tree\_Processing* program and the property *Prop* into a set of CHCs that are satisfiable iff *Prop* holds for *Tree\_Processing*. We get the following set of clauses (clauses 1–7 for the program and clause 8 for the property):

- 1.  $min(X, Y, Z) \leftarrow X < Y, Z = X$
- 2.  $min(X, Y, Z) \leftarrow X \ge Y, Z = Y$
- 3.  $min\_leaf(leaf, M) \leftarrow M = 0$

4.  $min\_leaf(node(X, L, R), M) \leftarrow M = M3+1, min\_leaf(L, M1), min\_leaf(R, M2), min(M1, M2, M3)$ 

- 5.  $left\_drop(N, leaf, leaf) \leftarrow$
- 6.  $left_drop(N, node(X, L, R), node(X, L, R)) \leftarrow N \leq 0$
- 7.  $left_drop(N, node(X, L, R), T) \leftarrow N \ge 1, N1 = N-1, left_drop(N1, L, T)$

8. false  $\leftarrow N \ge 0$ , M + N < K, left\_drop(N, T, U), min\_leaf(U, M), min\_leaf(T, K)

In clause 8 the head is *false* because the relation '<' in the constraint M+N < K occurring in the body is the negation of the relation ' $\geq$ ' occurring in the property *Prop*.

The Z3 solver is *not* able to prove the satisfiability of clauses 1-8 (which amounts to show that *false* is not derivable), because of the presence of variables of type *tree*. One can extend the capabilities of Z3 by adding induction rules on trees as done in recent work (Reynolds and Kuncak 2015, Suter et al. 2011, Unno et al. 2017). Instead, as we propose in this paper, we can derive by transformation, starting from clauses 1-8, a new, equisatisfiable set of clauses without variables of type *tree* and whose constraints are in *LIA* only, and then we can use CHC solvers, such as Z3, to show the satisfiability of this new set of clauses. Starting from clauses 1-8, our transformation algorithm, whose details are given in Section 4, produces the following clauses, where *min* is defined by clauses 1 and 2:

 $\begin{array}{l} 9. \ new1(N,M,K) \leftarrow M = 0, \ K = 0 \\ 10. \ new1(N,M,K) \leftarrow N \leq 0, \ M = M3 + 1, \ K = M, \ new2(M1), \ new2(M2), \\ min(M1,M2,M3) \\ 11. \ new1(N,M,K) \leftarrow N \geq 1, N1 = N - 1, K = K3 + 1, \ new1(N1,M,K1), \ new2(K2), \\ min(K1,K2,K3) \end{array}$ 

- 12.  $new2(M) \leftarrow M = 0$
- 13.  $new2(M) \leftarrow M = M3 + 1$ , new2(M1), new2(M2), min(M1, M2, M3)
- 14.  $false \leftarrow N \ge 0, M+N < K, new1(N, M, K)$

Now, the Z3 solver for CHCs with LIA constraints is able to prove the satisfiability of clauses 1, 2, 9–14 without using any induction on trees. The details of the transformation from clauses 1–8 to clauses 1, 2, 9–14 will be presented in Section 4.

## **3** Preliminaries

Let us consider a typed, first order functional language. We assume that the *basic types* of the language are *int*, for integers, and *bool*, for booleans. We also have *non-basic types*, which are introduced by (possibly recursive) type definitions such as the one for trees of integers considered in Section 2. The type system of our language can be formally defined as follows. (One may consider richer type systems including, for instance, parameterized types, but this system is sufficient for presenting our verification technique.)

 $Types \ni \tau ::= int \mid bool \mid ident \mid \tau_1 * \dots * \tau_k$  $Type \ Definitions: ident = c_1 \text{ of } \tau_1 \mid \dots \mid c_n \text{ of } \tau_n$ 

where: (i) *ident* is a type identifier, (ii) the operator '\*' builds k-tuple types, (iii)  $c_1, \ldots, c_n$  are distinct constructors with arity, and (iv) in any expression ' $c_i$  of  $\tau_i$ ', if  $c_i$  has arity k > 0, then  $\tau_i$  is a k-tuple type, and if  $c_i$  has arity 0, then 'of  $\tau_i$ ' is absent.

Every function has a *functional type* of the form  $\tau_1 \to \tau_2$ , for some types  $\tau_1$  and  $\tau_2$ , modulo the usual isomorphism between  $\tau \to (\tau' \to \tau'')$  and  $(\tau * \tau') \to \tau''$ .

In the following definitions: (i) n is an integer, (ii) b is *true* or *false*, (iii) x is a typed variable, (iv) c is a constructor of arity  $k (\geq 0)$ , and (v) f is a function of arity  $k (\geq 0)$  or a primitive operator on integers or booleans such as:  $+, \times, =, \neq, \leq, \neg, \land$ , and  $\Rightarrow$ .

We assume that all values, terms, and patterns are well-typed. In every let-in term x is a new variable not occurring in  $t_0$  and occurring in  $t_1$ . In every match-with term the patterns  $p_1, \ldots, p_k$  are pairwise disjoint and exhaustive. A user-defined function f is defined by: let rec  $f x_1 \ldots x_k = t$ , where the free variables of t are among  $\{x_1, \ldots, x_k\}$ .

A program is a (possibly empty) set of type definitions together with a set of user-defined functions. Given a program P and a term t without free variables and whose functions are defined in P, the value v of t using P is computed according to the call-by-value semantics. In this case we write  $t \to_P v$ . We say that a program P terminates if, for every term t without free variables and whose functions are defined in P, there exists a value vsuch that  $t \to_P v$ . Given a boolean term q whose free variables are in  $\{x_1, \ldots, x_n\}$ , a property is a universally quantified formula of the form  $\forall x_1, \ldots, x_n.q$ . Given a program P, we say that a property  $\forall x_1, \ldots, x_n.q$ , whose functions are defined in P, holds for P iff for all values  $v_1, \ldots, v_n$ , we have that  $q[v_1, \ldots, v_n / x_1, \ldots, x_n] \to_P true$ , where the square bracket notation is for substitution.

Let us consider a typed first order logic (Enderton 1972) which includes: (i) the theory

LIA of the linear integer arithmetic constraints, and (ii) the theory Bool of boolean constraints. A constraint in  $LIA \cup Bool$  is any formula in  $LIA \cup Bool$ .

An *atom* is a formula of the form  $q(t_1, \ldots, t_m)$ , where q is a typed predicate symbol not used in  $LIA \cup Bool$ , and  $t_1, \ldots, t_m$  are typed terms made out of variables and constructors. A *constrained Horn clause* (or simply, a *clause*, or a CHC) is an implication of the form  $A \leftarrow c, B$  (comma denotes conjunction), where the conclusion (or *head*) A is either an atom or *false*, the premise (or *body*) is the conjunction of a constraint c, and a (possibly empty) conjunction B of atoms. A clause whose head is an atom is called a *definite clause*, and a clause whose head is *false* is called a *goal*. A *constrained fact* is a clause of the form  $A \leftarrow c$ . We will write the constrained fact  $A \leftarrow true$  also as  $A \leftarrow$ . We assume that all variables in a clause are universally quantified in front. Given a term t, by vars(t)we denote the set of variables occurring in t. Similarly for the set of variables occurring in atoms, or clauses, or sets of atoms, or sets of clauses. A set S of CHCs is said to be *satisfiable* if  $S \cup LIA \cup Bool$  has a model, or equivalently,  $S \cup LIA \cup Bool \not\models false$ .

We define a translation Tr from any set P of function definitions into a set of CHCs by: (i) introducing for each function f of arity k, a new predicate, say p, of arity k+1, and then (ii) providing the clauses for predicate p so that  $f(x_1, \ldots, x_k) = y$  iff  $p(x_1, \ldots, x_k, y)$  holds. This translation has been applied in Section 2 for generating clauses 1–7 starting from the *Tree\_Processing* program. Primitive operations, such as + and -, are translated in terms of constraints expressed in *LIA*. Similarly, a property of the form:  $\forall x. r(x) \Rightarrow s(x)$ is translated into a goal (or a set of goals, in general) as indicated in Section 2 (see goal 8). If we denote by Tr(P) and Tr(Prop) the set of CHCs generated from a given program P and a given property *Prop*, respectively, we have the following theorem, where satisfiability is defined within the typed logic we consider.

# $Theorem \ 1$

Given a program P that terminates, a property Prop holds for P iff the set  $Tr(P) \cup Tr(Prop)$  of clauses is satisfiable.

## 4 Eliminating Inductively Defined Data Structures

Now we present an algorithm, called Algorithm  $\mathcal{E}$ , for eliminating from CHCs the predicate arguments that have a non-basic type, such as lists or trees. If Algorithm  $\mathcal{E}$  terminates, then it transforms a set of clauses into an equisatisfiable set, where the arguments of all predicates have basic types. The algorithm is based on the fold/unfold strategy for eliminating unnecessary variables from logic programs (Proietti and Pettorossi 1995).

Given two terms  $t_1$  and  $t_2$ , by  $t_1 \leq t_2$  and  $t_1 \prec t_2$  we denote the *subterm* and *strict* subterm relation, respectively. Given an atom A, by nbargs(A) we denote the set of arguments of non-basic type of A. By nbvars(A) we denote the set of variables of non-basic type occurring in A. The next definition extends the  $\leq$  and  $\prec$  relations to atoms.

## Definition 1 (Atom Comparison)

Given two atoms  $A_1$  and  $A_2$ ,  $A_1 \leq A_2$  (or  $A_1 \prec A_2$ ) holds if there exist  $t_1 \in nbargs(A_1)$ and  $t_2 \in nbargs(A_2)$  such that: (i)  $vars(t_1) \cap vars(t_2) \neq \emptyset$ , and (ii)  $t_1 \leq t_2$  (or  $t_1 \prec t_2$ ). Given a set S of atoms, an atom  $M \in S$  is strictly maximal in S if there exists no atom  $A \in S$ such that  $M \prec A$  and there exists an atom  $A' \in S$  such that  $A' \prec M$ . For example, q([Y|Ys], []) is strictly maximal in the set  $\{p([X], Ys), q([Y|Ys], [])\}$ . Indeed,  $Ys \prec [Y|Ys]$ , and hence  $p([X], Ys) \prec q([Y|Ys], [])$ , while  $q([Y|Ys], []) \not\prec p([X], Ys)$ (note that  $[] \prec [X]$ , but  $vars([]) \cap vars([X]) = \emptyset$ ). The notion of a strictly maximal atom will be used in the Unfold procedure of Algorithm  $\mathcal{E}$  to guide the unfolding process.

A predicate *has basic types* if *all* its arguments have basic type. An atom has basic types if its predicate has basic types. A clause has basic types if *all* its atoms have basic types.

## Definition 2 (Sharing Blocks)

Let S be a set (or a conjunction) of atoms. For any two atoms  $A_1$  and  $A_2$  in S,  $A_1 \downarrow A_2$ holds iff  $nbvars(A_1) \cap nbvars(A_2) \neq \emptyset$ . Let  $\Downarrow$  be the reflexive transitive closure of  $\downarrow$ . By *SharingBlocks*(S) we denote the partition of S into subsets, called *sharing blocks*, with respect to  $\Downarrow$ .

Algorithm  $\mathcal{E}$  makes use of the well-known transformation rules *define*, *fold*, *unfold*, and *replace* for CHCs (Etalle and Gabbrielli 1996, Tamaki and Sato 1984).

# The Elimination Algorithm $\mathcal{E}$ .

Input: A set  $Cls \cup Gs$ , where Cls is a set of definite clauses and Gs is a set of goals; *Output*: A set *TransfCls* of clauses such that: (1)  $Cls \cup Gs$  is satisfiable iff *TransfCls* is satisfiable, and (2) every clause in *TransfCls* has basic types.

 $\begin{array}{l} Defs := \emptyset; \ InCls := Gs; \ TransfCls := \emptyset; \\ \textbf{while} \ InCls \neq \emptyset \ \textbf{do} \\ Define-Fold(Defs, InCls, NewDefs, FldCls); \\ Unfold(NewDefs, Cls, UnfCls); \\ Replace(UnfCls, Cls, RCls); \\ Defs := Defs \cup NewDefs; \ InCls := RCls; \ TransfCls := TransfCls \cup FldCls; \\ \end{array}$ 

Starting from a set *Cls* of definite clauses and a set *Gs* of goals, Algorithm  $\mathcal{E}$  applies iteratively the procedures *Define-Fold*, *Unfold*, and *Replace*, in this order, until it derives a set *TransfCls* of clauses whose predicates have basic types only. Algorithm  $\mathcal{E}$  collects in *Defs* the clauses, called *definitions*, introduced by the *Define* steps, and collects in *InCls* the clauses to be transformed. The set *Defs* of definitions is initialized to the empty set and the set *InCls* of the input goals is initialized to the set *Gs* of goals.

Now let us present the various procedures used by Algorithm  $\mathcal{E}$ .

 $\begin{array}{l} \textbf{Procedure } Define-Fold(Defs, InCls, NewDefs, FldCls)\\ Input: A set Defs of definitions and a set InCls of clauses;\\ \underline{Output: A set NewDefs of definitions and a set FldCls of clauses.}\\ \hline NewDefs := \emptyset; \ FldCls := \emptyset;\\ \textbf{for each clause } C: \ H \leftarrow c, B \ \text{in } InCls \ \textbf{do}\\ \textbf{if } C \ \text{is a constrained fact } \textbf{then } FldCls := FldCls \cup \{C\} \ \textbf{else}\\ \underline{Define. \ Let \ SharingBlocks(B) = \{B_1, \dots, B_n\};}\\ \textbf{for } i = 1, \dots, n \ \textbf{do}\\ \textbf{if there is no clause in } Defs \cup NewDefs \ \textbf{whose body is } B_i \ (\text{modulo the names of variables and the order and multiplicity of the atoms) } \textbf{then} \end{array}$ 

 $NewDefs := NewDefs \cup \{newp_i(V_i) \leftarrow B_i\}$ where: (i)  $newp_i$  is a new predicate symbol, and (ii)  $V_i$  is the tuple of distinct variables of basic type occurring in  $B_i$ ; Fold. C is folded using the definitions in  $Defs \cup NewDefs$ , thereby deriving  $\overline{F}: H \leftarrow c, newp_1(V_1), \dots, newp_n(V_n)$ where, for  $i = 1, \dots, n$ ,  $newp_i(V_i) \leftarrow B_i$  is the unique clause in  $Defs \cup NewDefs$ whose body is  $B_i$ , modulo variable renaming;  $FldCls := FldCls \cup \{F\}$ ;

The Define-Fold procedure removes the arguments with non-basic types from each clause C of InCls in two steps: first, the Define step introduces a new predicate definition for each sharing block of the body of C (unless a definition with a body equal, modulo variable renaming, to that block has already been introduced in a previous Define step), and then, the Fold step replaces each sharing block by the head of the new definition. Since the heads of these new definitions have basic types, the body of the clause of the form:  $H \leftarrow c, newp_1(V_1), \ldots, newp_n(V_n)$ , derived from C by the Fold step, has basic types. Also H has basic types, because: (i) Algorithm  $\mathcal{E}$  initializes InCls to the set Gs of goals, whose heads are false, and (ii) the head predicate of each new clause added to InCls has been introduced by a previous Define step, and hence, by construction, has basic types.

## Example 1

(*Tree\_Processing*). Let us consider the introductory example of Section 2. At the first iteration of the while-do body of Algorithm  $\mathcal{E}$ , *InCls* consists of clause 8, which is the result of translating the property *Prop* to be verified. The *Define* step introduces the new predicate *new*1 through the following definition:

15.  $new1(N, M, K) \leftarrow left\_drop(N, T, U), min\_leaf(U, M), min\_leaf(T, K)$ 

The body of this clause consists of the single sharing block of the body of clause 8 (note that  $left\_drop(N,T,U)$  shares the tree variables T and U with  $min\_leaf(T,K)$  and  $min\_leaf(U,M)$ , respectively). Now, the Fold step derives clause 14 of Section 2, where all predicates have arguments of integer type, and thus it is added to the final set TransfCls.

The *Define* step adds to *Defs* a set *NewDefs* of new definitions, whose body may have predicates with non-basic types (see clause 15). Now, Algorithm  $\mathcal{E}$  proceeds by applying the *Unfold* procedure to those clauses.

# **Procedure** Unfold(NewDefs, Cls, UnfCls)

*Input*: A set *NewDefs* of definitions and a set *Cls* of definite clauses; *Output*: A set *UnfCls* of clauses.

UnfCls := NewDefs;

Initially, all atoms in the body of the clauses of UnfCls are marked 'unfoldable';

while there exists a clause C in UnfCls of the form  $H \leftarrow c, L, A, R$  such that either

(i) the atom A is 'unfoldable' and is strictly maximal in L, A, R, or

(ii) all atoms in L, A, R are 'unfoldable' and not strictly maximal do

Let  $K_1 \leftarrow c_1, B_1, \ldots, K_m \leftarrow c_m, B_m$  be all clauses of Cls (where, without loss of generality, we assume  $vars(Cls) \cap vars(C) = \emptyset$ ) such that, for  $i = 1, \ldots, m$ , (1) there exists a most general unifier  $\vartheta_i$  of A and  $K_i$ , and (2) the constraint  $(c, c_i)\vartheta_i$  is satisfiable;

 $UnfCls := (UnfCls - \{C\}) \cup \{(H \leftarrow c, c_1, L, B_1, R)\vartheta_1, \dots, (H \leftarrow c, c_m, L, B_m, R)\vartheta_m\}$ where, for  $i = 1, \dots, m$ , (1) an atom  $E\vartheta_i$  of  $(L, R)\vartheta_i$  is 'unfoldable' iff the corresponding atom E of (L, R) is 'unfoldable' in C, and (2) no atom in  $B\vartheta_i$  is 'unfoldable';

The Unfold procedure unfolds the atoms occurring in NewDefs by performing resolution steps with clauses in Cls. The procedure applies a strategy that consists in unfolding strictly maximal atoms, if any. In the case where the predicates are defined by induction on the structure of their arguments with non-basic types, this strategy corresponds to a form of induction on the arguments structure. The subsequent folding steps correspond to applications of the inductive hypotheses. A characterization of a class of CHCs where this strategy is successful will be given in Section 5. The use of the 'unfoldable' marking on atoms enforces a finite number of resolution steps.

Note that when Case (ii) of the condition of the while-do holds, the *Unfold* procedure may unfold *any* 'unfoldable' atom (and in our implementation of the procedure we unfold the leftmost one). However, in the class of CHCs presented in Section 5, the termination of Algorithm  $\mathcal{E}$  is independent of the choice of the atom to be unfolded.

## Example 2

(*Tree\_Processing, continued*). Let us continue the derivation presented in Example 1. All atoms in the body of clause 15 are 'unfoldable' and none is strictly maximal (indeed, no tree argument is a strict superterm of another). In this case the *Unfold* procedure may unfold any 'unfoldable' atom, and we assume that it unfolds  $left_drop(N,T,U)$ . The following three clauses, where we have underlined the 'unfoldable' atoms, are derived:

 $\begin{array}{ll} 16. \ new1(N,M,K) \leftarrow \underline{min\_leaf(leaf,M)}, \ \underline{min\_leaf(leaf,K)} \\ 17. \ new1(N,M,K) \leftarrow \overline{N \leq 0}, \ \underline{min\_leaf(node(X,L,R),M)}, \ \underline{min\_leaf(node(X,L,R),K)} \\ 18. \ new1(N,M,K) \leftarrow N \geq 1, \ \overline{N1=N-1}, \ left\_drop(N1,L,U), \ \underline{min\_leaf(U,M)}, \\ min\_leaf(node(X,L,R),K) \end{array}$ 

Now the Unfold procedure continues by selecting more atoms for unfolding. For instance, in clause 18 it selects  $min\_leaf(node(X,L,R),K)$ , which is strictly maximal because the argument L of  $left\_drop(N1,L,U)$  is a strict subterm of node(X,L,R) and no atom in the body of clause 18 has an argument that is a strict superterm of node(X,L,R). After five iterations, where all underlined atoms, except  $min\_leaf(U,M)$ , are unfolded, the Unfold procedure derives the following clauses:

- 19.  $new1(N, M, K) \leftarrow M = 0, K = 0$
- $\begin{array}{l} 20. \ new1(N,M,K) \leftarrow N \leq 0, \ M = M3 + 1, \ K = K3 + 1, \ min\_leaf(L,M1), \ min\_leaf(R,M2), \\ min(M1,M2,M3), \ min\_leaf(L,K1), \ min\_leaf(R,K2), \ min(K1,K2,K3) \end{array}$
- 21.  $new1(N, M, K) \leftarrow N \ge 1$ , N1 = N-1 K = K3+1,  $left\_drop(N1, L, U)$ ,  $\underline{min\_leaf(U, M)}$ ,  $min\_leaf(L, K1)$ ,  $min\_leaf(R, K2)$ , min(K1, K2, K3)

In clause 21,  $min\_leaf(U, M)$  is an 'unfoldable' atom, but the Unfold procedure stops because that atom is not strictly maximal. Clause 19 is a constrained fact and it is added to the final set TransfCls (indeed, clause 19 is clause 9 of Section 2).

After the Unfold procedure Algorithm  $\mathcal{E}$  may simplify some clauses by exploiting functional dependencies among predicate arguments, which hold by construction for the predicates obtained by translating functional programs.

## Definition 3 (Predicate Functionality)

Let *Cls* be a set of definite clauses. A predicate p(X,Y), where X and Y are tuples of arguments, is *functional* in *Cls* if  $Cls \cup \{false \leftarrow Y \neq Z, p(X,Y), p(X,Z)\}$  is a satisfiable set of clauses.

## **Procedure** Replace(UnfCls, Cls, RCls)

Input: A set UnfCls of clauses and a set Cls of definite clauses;

*Output*: A set *RCls* of clauses.

## RCls := UnfCls;

while there is a clause  $C \in RCls$  of the form:  $H \leftarrow c, G_1, p(t, u), G_2, p(t, w), G_3$ , where predicate p(X,Y) is functional in Cls do

Replace C by  $(H \leftarrow c, G_1, p(t, u), G_2, G_3)\vartheta$ , where  $\vartheta$  is a most general unifier of u and w;

# Example 3

(*Tree\_Processing, continued*). By the functionality of the predicates  $min\_leaf(T, M)$  and min(X, Y, Z), clause 20 is replaced by the following one:

22.  $new1(N,M,K) \leftarrow N \leq 0, M = M3+1, K = M, min\_leaf(L,M1), min\_leaf(R,M2), min(M1,M2,M3)$ 

Now, Algorithm  $\mathcal{E}$  performs a second iteration by executing again the *Define-Fold*, *Unfold*, and *Replace* procedures. The *Define* step introduces the following definition:

23.  $new2(M1) \leftarrow min\_leaf(L, M1)$ 

whose body consists of a sharing block in the body of clause 22 (note that the tree variable L is not shared with any other atom). No other definitions are introduced, as all other sharing blocks in the clauses currently in *InCls* (namely, clauses 19, 21, and 22) are variants of the body of the definitions 15 and 23. The *Fold* step derives clauses 10 and 11 of Section 2 from clauses 22 and 21, respectively.

Finally, Algorithm  $\mathcal{E}$  performs an Unfold step followed by a Fold step on clause 23 and derives clauses 12 and 13 of Section 2. Note that Replace steps are not applicable, and no new definitions are introduced by the Define step. Thus, Algorithm  $\mathcal{E}$  terminates and returns clauses 1, 2, 9–14 of Section 2.

## 5 Correctness and Termination of the Transformation

The partial correctness of Algorithm  $\mathcal{E}$  follows from well-known satisfiability preservation results that hold for the fold/unfold transformation rules (Etalle and Gabbrielli 1996, Tamaki and Sato 1984) (see the Appendix for a proof).

# Theorem 2 (Partial Correctness)

Let Cls be a set of definite clauses and let Gs be a set of goals. If Algorithm  $\mathcal{E}$  terminates for the input clauses  $Cls \cup Gs$ , returning a set TransfCls of clauses, then (1)  $Cls \cup Gs$  is satisfiable iff TransfCls is satisfiable, and (2) all clauses in TransfCls have basic types.

Now we introduce a class of CHCs where Algorithm  $\mathcal{E}$  is guaranteed to terminate.

First we need the following terminology and notation. An atom is said to be *linear* if each variable occurs in it at most once. By arity(p) we denote the arity of predicate symbol p.

## Definition 4 (Slice Decomposition)

Let C be a clause of the form  $A_0 \leftarrow c, A_1, \ldots, A_m$ , and let *Pred* be the set of predicate symbols in the atoms  $A_0, A_1, \ldots, A_m$  of C. A *slice* of C is a total function  $\sigma : Pred \to \mathbb{N}$ such that, for every predicate  $p \in Pred$ : (1)  $0 \le \sigma(p) \le arity(p)$ , and (2) if  $\sigma(p) = i > 0$ , then the *i*-th argument of p has a non-basic type. For any atom  $p(t_1, \ldots, t_k)$  in C, we define:

$$\sigma(p(t_1,\ldots,t_k)) = \begin{cases} p_0 & \text{if } \sigma(p) = 0\\ p_i(t_i) & \text{if } \sigma(p) = i > 0 \end{cases}$$

where  $p_0$  (of arity 0) and  $p_i$  (of arity 1) are fresh, new predicate symbols.

We also define  $\sigma(C)$  to be the new clause  $\sigma(A_0) \leftarrow \sigma(A_1), \ldots, \sigma(A_m)$ . The slice  $\sigma$  of C is quasi-descending if  $\sigma(A_0), \sigma(A_1), \ldots, \sigma(A_m)$  are linear atoms and, for  $i = 1, \ldots, m$ , (1) for  $j = 1, \ldots, m$ ,  $vars(\sigma(A_i)) \cap vars(\sigma(A_j)) = \emptyset$ , if  $i \neq j$ , and (2) either  $vars(\sigma(A_0)) \cap vars(\sigma(A_i)) = \emptyset$  or  $\sigma(A_0) \succeq \sigma(A_i)$ .

A slice decomposition of clause C is a set  $\Sigma_C = \{\sigma_1, \ldots, \sigma_n\}$ , where  $\sigma_1, \ldots, \sigma_n$  are slices of C and, for all  $p \in Pred$ , for all  $i \in \{1, \ldots, arity(p)\}$ , if the *i*-th argument of p has nonbasic type, then there exists  $\sigma_j \in \Sigma_C$  such that  $\sigma_j(p) = i$ . A slice decomposition  $\Sigma_C$  of a clause C is said to be *disjoint* if, for any two slices  $\sigma_h$  and  $\sigma_k$  in  $\Sigma_C$ , we have that  $vars(\sigma_h(C)) \cap vars(\sigma_k(C)) = \emptyset$ , whenever  $h \neq k$ .  $\Sigma_C$  is said to be *quasi-descending* if all slices in  $\Sigma_C$  are quasi-descending.

The following are slices of clauses 4 and 7 (see Section 2):

 $\sigma_2(\text{clause } 7) = left_drop_2(node(X, L, R)) \leftarrow left_drop_2(L)$ 

 $\sigma_3(\text{clause 7}) = left\_drop_3(T) \leftarrow left\_drop_3(T)$ 

We have that  $\Sigma_4 = \{\sigma_1\}$  is a disjoint, quasi-descending slice decomposition of clause 4, and so is  $\Sigma_7 = \{\sigma_2, \sigma_3\}$  for clause 7.

Now we define a class of goals for which Algorithm  $\mathcal{E}$  terminates.

## Definition 5 (Sharing Cycle)

Let G be a goal of the form  $false \leftarrow c, A_1, \ldots, A_m$ . We say that G has a sharing cycle if there is a sequence of k (> 1) distinct variables  $X_0, \ldots, X_{k-1}$  of non-basic type, and a sequence  $A'_0, \ldots, A'_k$  of atoms in  $A_1, \ldots, A_m$ , such that: (i)  $A'_0 = A'_k$ , (ii)  $A'_1, \ldots, A'_k$  are all distinct, and (iii) for  $i = 0, \ldots, k-1, A'_i$  and  $A'_{i+1}$  share the non-basic variable  $X_i$ .

Now, if we represent the body of goal 8 of Section 2 as the following labeled graph:

$$min\_leaf(T,K) \xrightarrow{T} left\_drop(N,T,U) \xrightarrow{U} min\_leaf(U,M)$$

where the arc from a node A to a node B has label X iff  $X \in nbvars(A) \cap nbvars(B)$ , then it is easy to see that goal 8 has no sharing cycle (indeed, there are no cycles in the above graph).

We have the following result, whose proof is given in the Appendix.

## Theorem 3 (Termination)

Let Cls be a set of definite clauses such that every clause in Cls has a disjoint, quasidescending slice decomposition. Let Gs be a set of goals such that, for each goal  $G \in Gs$ , (i) G is of the form  $false \leftarrow c, A_1, \ldots, A_m$ , where for  $i = 1, \ldots, m$ ,  $A_i$  is an atom whose arguments are distinct variables, and (ii) G has no sharing cycles. Then Algorithm  $\mathcal{E}$ terminates for the input clauses  $Cls \cup Gs$ . Note that all definite clauses (clauses 1-7) of our introductory example have a quasidescending slice decomposition (see above for clauses 4 and 7). These slice decompositions are also disjoint, except for the one of clause 6 where the second and third argument of *left\_drop* share some variables. However, clause 6 can be rewritten as:

 $\begin{array}{l} \textit{left\_drop}(N,\textit{node}(X,L,R),\textit{node}(X1,L1,R1)) \leftarrow N \leq 0, \ X = X1, \ \textit{eqt}(L,L1), \ \textit{eqt}(R,R1) \\ \textit{eqt}(\textit{leaf},\textit{leaf}) \leftarrow \end{array}$ 

 $eqt(node(X1,L1,R1),node(X2,L2,R2)) \leftarrow X1 = X2, \; eqt(L1,L2), \; eqt(R1,R2)$ 

where predicate eqt defines the equality between binary trees. These three clauses have a disjoint, quasi-descending slice decomposition, and after this rewriting the termination of Algorithm  $\mathcal{E}$  is guaranteed. The rewriting of any constrained fact, such as clause 6, into a clause that has a disjoint, quasi-descending slice decomposition can be done automatically as a pre-processing step, by introducing an equality predicate for each non-basic type. However, in the benchmark set presented in Section 7, this pre-processing step has no effect on the termination behavior of our transformation algorithms.

# 6 Adding Integer and Boolean Constraints

Algorithm  $\mathcal{E}$  is not guaranteed to terminate outside the class of definite clauses and goals considered in Theorem 3. Let us consider, for instance, the following set of clauses which specifies a verification problem on lists:

| $append([], Ys, Ys) \leftarrow$                        | $take(N, [], []) \leftarrow$                            |
|--|---|
| $append([X Xs], Ys, [Z Zs]) \leftarrow X = Z,$         | $take(N, [X Xs], []) \leftarrow N = 0$                  |
| append(Xs, Ys, Zs)                                     | $take(N, [X Xs], [Y Ys]) \leftarrow N \neq 0, \ X = Y,$ |
|  | N1 = N - 1, take(N1, Xs, Ys)                            |
| $drop(N, [], []) \leftarrow$                           | $diff\_list([],[Y Ys]) \leftarrow$                      |
| $drop(N, [X Xs], [Y Xs]) \leftarrow N = 0, X = Y$      | $diff\_list([X Xs],[]) \leftarrow$                      |
| $drop(N, [X Xs], Ys) \leftarrow N \neq 0, N1 = N - 1,$ | $diff\_list([X Xs], [Y Ys]) \leftarrow X \neq Y$        |
| drop(N1, Xs, Ys)                                       | $diff\_list([X Xs], [Y Ys]) \leftarrow X = Y,$          |
|  | $diff\_list(Xs, Ys)$                                    |

 $false \leftarrow M = N, take(M, Xs, Ys), drop(N, Xs, Zs), append(Ys, Zs, A), diff_list(A, Xs)$ 

In these clauses: (i) take(M, Xs, Ys) holds if the list Ys is the prefix of the list Xs up to its M-th element, (ii) drop(N, Xs, Zs) holds if list Zs is a suffix of Xs starting from its (N+1)-th element, (iii) append(Ys, Zs, A) holds if A is the concatenation of the lists Ysand Zs, and (iv)  $diff_{list}(A, Xs)$  holds if A and Xs are different lists.

The definite clauses listed above satisfy the hypothesis of Theorem 3 (after rewriting some constrained facts as done at the end of the previous section), but the goal does not. Indeed, that goal has the following sharing cycle:

take(M, Xs, Ys) <u>Xs</u> drop(N, Xs, Zs) <u>Zs</u> append(Ys, Zs, A) <u>Ys</u> take(M, Xs, Ys)Algorithm  $\mathcal{E}$  does not terminate on this example. Indeed, starting from the definition:

 $new0(M, N) \leftarrow take(M, Xs, Ys), drop(N, Xs, Zs), append(Ys, Zs, A), diff_list(A, Xs)$ 

infinitely many new predicates with unbounded lists are generated by Algorithm  $\mathcal{E}$ . However, these predicates correspond to cases where  $M \neq N$ , and if we keep the constraint M=N between the first arguments of *take* and *drop*, then a finite set of new predicates is generated. In particular, if we start from the definition:  $new1(M, N) \leftarrow M=N, take(M, Xs, Ys), drop(N, Xs, Zs), append(Ys, Zs, A), diff_list(A, Xs)$ the transformation terminates after a few steps and derives the following equisatisfiable set of clauses, where an equality constraint holds for the two arguments of each occurrence

 $\begin{array}{l} new2 \leftarrow new2\\ new1(M,N) \leftarrow M=N, \ M=0, \ new2\\ new1(M,N) \leftarrow M=N, \ M\neq0, \ M=1+M1, \ N=1+N1, \ new1(M1,N1)\\ false \leftarrow M=N, \ new1(M,N) \end{array}$ 

The satisfiability of this set of clauses is trivial, because it does not contain any constrained fact, and is easily proved by the Z3 solver.

The above example motivates the introduction of a variant of Algorithm  $\mathcal{E}$ , called Algorithm  $\mathcal{EC}$ , which is obtained by allowing in the *Define* step the introduction of definitions whose bodies may include constraints in  $LIA \cup Bool$ , and hence applying *Fold* steps with respect to these definitions. The rest of Algorithm  $\mathcal{EC}$  is equal to Algorithm  $\mathcal{E}$ .

As usual in constraint-based transformation techniques (see (De Angelis et al. 2017a) for a recent paper), the computation of a suitable constraint when introducing a new definition is done by means of a constraint generalization function Gen. We say that a constraint g is more general than, or it is a generalization of, a constraint c, if  $LIA \cup Bool \models$  $\forall (c \rightarrow g)$ . Given a constraint c, a conjunction B of atoms, and a set Defs of definitions, the function Gen matches the constraint c against the constraint d occurring in a definition in Defs whose body is of the form d, B (modulo variable renaming), and returns a new constraint Gen(c, B, Defs) which is more general than both c and d.

The details of how the constraint generalization function Gen is actually implemented are not necessary for understanding Algorithm  $\mathcal{EC}$ , which is parametric with respect to such a function. Let us only mention here that the function Gen used for the experiments reported in Section 7 makes use on the *widening operator* based on *bounded difference shapes*, which is a standard operator on convex polyhedra considered in the field of *abstract interpretation* (Bagnara et al. 2008, Cousot and Halbwachs 1978). The use of widening avoids the introduction of infinitely many definitions that differ for the constraints only.

The *Define* and *Fold* steps for Algorithm  $\mathcal{EC}$  are as follows.

Let  $C: H \leftarrow c, B$  in *InCls* be a clause which is not a constrained fact.

<u>Define</u>. Let  $SharingBlocks(B) = \{B_1, \ldots, B_n\};$ 

for  $i = 1, \ldots, n$  do

 $g_i := Gen(c, B_i, Defs \cup NewDefs);$ 

if there is no clause in  $Defs \cup NewDefs$  whose body is  $g_i, B_i$ 

then  $NewDefs := NewDefs \cup \{newp_i(V_i) \leftarrow g_i, B_i\}$ 

where: (i)  $newp_i$  is a new predicate symbol, and (ii)  $V_i$  is the tuple of distinct variables of basic type occurring in  $B_i$ ;

Fold. C is folded using the definitions in  $Defs \cup NewDefs$ , thereby deriving

 $F: H \leftarrow c, newp_1(V_1), \dots, newp_n(V_n)$ 

where, for i = 1, ..., n,  $newp_i(V_i) \leftarrow g_i, B_i$  is the unique clause in  $Defs \cup NewDefs$ whose body is  $g_i, B_i$ , modulo variable renaming;

 $FldCls := FldCls \cup \{F\};$ 

of new1:

The partial correctness of Algorithm  $\mathcal{E}$  (see Theorem 2) carries over to Algorithm  $\mathcal{EC}$ , because also  $\mathcal{EC}$  can be expressed as a sequence of applications of the fold/unfold transformation rules for CHCs (Etalle and Gabbrielli 1996). Also the termination result for Algorithm  $\mathcal{E}$  (see Theorem 3) extends to  $\mathcal{EC}$ , as long as the function Gen guarantees that, for a given conjunction B of atoms, finitely many definitions of the form  $newp(V) \leftarrow g, B$ can be introduced.

#### 7 Experimental Evaluation

In this section we present the results of an experimental evaluation we have performed for assessing the effectiveness of our approach and, in particular, for comparing it with the approach that extends CHC solvers by adding inductive rules, as done in the RCAML tool (Unno et al. 2017) based on the Z3 solver.

*Implementation.* We have implemented the transformation strategy presented in Section 4 using the VERIMAP system (De Angelis et al. 2014) together with the Parma Polyhedra Library (PPL) (Bagnara et al. 2008) for performing constraint generalizations. Then, we have used the Z3 solver v4.6.0 with the SPACER fixed-point engine (Komuravelli et al. 2013) to check the satisfiability of the transformed CHCs.

The tool and the benchmark suite are available at https://fmlab.unich.it/iclp2018/.

Benchmark suite and experiments. Our benchmark suite is a collection of 105 verification problems, each one consisting of an OCaml functional program manipulating inductively defined data structures (such as lists or trees) together with a property to be verified. Most of the problems (70 out of 105) derive from the benchmark suite of RCAML (see http://www.cs.tsukuba.ac.jp/~uhiro/, Software RCaml, [web demo (induction)]). This suite, in turn, includes problems from the suite of IsaPlanner (Dixon and Fleuriot 2003) which is a generic framework for proof planning in the Isabelle theorem prover. We divide our benchmark suite into four sets of problems (see Table 1): (1) FirstOrder, the set of problems relative to first-order programs (57 out of 70) in the RCAML suite, (2) HigherOrderInstances, the set of problems relative to first-order programs (13 out of 70) that have been obtained by instantiating higher-order programs in the RCAML suite, (3) MoreLists, a set of 16 verification problems on lists, and (4) MoreTrees, a set of 19 verification problems on trees. In our benchmark 94 programs do satisfy the associated property and the remaining 11 do not.

By using the preprocessor provided by the RCAML system, each verification problem has been translated into a set of CHCs (see the translation Tr of Section 3). Then, for each derived set, call it I, of CHCs we have performed the following three experiments, whose results are summarized in Table 1. (A table with the detailed results for each problem of the benchmark is available at https://fmlab.unich.it/iclp2018/.)

- We have run Z3 (which does not use any structural induction rule) for checking the satisfiability of I (see the two columns for Z3).
- We have applied the transformation algorithm  $\mathcal{EC}$  to I, thereby producing a set T of CHCs, and then we have run Z3 for checking the satisfiability of T (see the two columns for  $\mathcal{EC}$ ;Z3).
- We have run RCAML on I (see the two columns for RCAML).

|                         |      | A COLORED AND A |              |                                |                                |                |                 |
|-------------------------|------|---|--------------|--------------------------------|--------------------------------|----------------|-----------------|
|                         |      | Z3  |              | $\mathcal{EC}; Z3$             |                                | RCAML          |                 |
| Problem Set             | n    | $S_{\rm Z3}$  | $T_{\rm Z3}$ | $S_{\mathcal{EC};\mathrm{Z3}}$ | $T_{\mathcal{EC};\mathrm{Z3}}$ | $S_{ m RCAML}$ | $T_{\rm RCAML}$ |
| (1) FirstOrder          | 57   | 3   | 0.09         | 47                             | 37.64                          | 41             | 216.59          |
| (2) HigherOrderInstance | s 13 | 1   | 0.04         | 11                             | 8.33                           | 10             | 45.40           |
| (3) MoreLists           | 16   | 3   | 13.87        | 14                             | 11.27                          | 10             | 119.01          |
| (4) More Trees          | 19   | 5   | 20.18        | 19                             | 26.79                          | 5              | 55.16           |
| Total                   | 105  | 12  | 34.18        | 91                             | 84.03                          | 66             | 436.17          |
| $Avg \ time$            |      |   | 2.85         |                                | 0.92                           |                | 6.61            |

Table 1. Column *n* reports the number of problems in each Problem Set. Columns  $S_{Z3}$  and  $T_{Z3}$  report the number of problems which have been solved by Z3 and the total time needed for their solution. Analogously, for the two columns referring to  $\mathcal{EC}$ ; Z3 and the two columns referring to RCAML. Times are in seconds. The timeout occurs after 300s.

For each verification problem we have set a timeout limit of 300 seconds. Z3 and VERIMAP have run on an Intel Xeon CPU E5-2640 2.00GHz with 64GB under CentOS. RCAML has run in a Linux virtual machine on an Intel is 2.3GHz with 8GB memory under macOS.

Results. The figures in Table 1 show that our approach considerably increases the effectiveness of CHC satisfiability checking. Indeed, when directly applied to the CHCs that encode the given verification problems, Z3 is able to solve 12 problems, while it solves 91 problems when it is applied to the CHCs produced by our transformation algorithm  $\mathcal{EC}$ (compare Columns  $S_{Z3}$  and  $S_{\mathcal{EC};Z3}$ ). For the remaining 14 problems, Algorithm  $\mathcal{EC}$  does not terminate within the time limit, and thus no CHCs are produced. Note that these 14 problems fall outside the class of programs for which Algorithm  $\mathcal{E}$ , and also Algorithm  $\mathcal{EC}$ , is guaranteed to terminate.

Table 1 shows that our approach compares quite well with respect to the induction based approach implemented in the RCAML system (Unno et al. 2017). Indeed,  $\mathcal{EC}$ ;Z3 proves 65 out of 66 problems that are also proved by RCAML, and in addition it proves 26 problems, 7 of which belong to the RCAML benchmark suite. Also the average times appear to be favorable to  $\mathcal{EC}$ ;Z3 with respect to RCAML. In particular, on the set of 67 problems where both  $\mathcal{EC}$ ;Z3 and RCAML provide the solution within the timeout,  $\mathcal{EC}$ ;Z3 is about six times faster than RCAML, having taken into account the fact that the machine on which we have run RCAML is approximately 13% slower.

Now let us report on some other important facts not shown in Table 1. When solving verification problems via  $\mathcal{EC}$ ; Z3, a substantial portion of work is performed by Algorithm  $\mathcal{EC}$ . Indeed, the total time spent by Z3 for solving the 91 problems is only 6% of the total solving time (84.03s). Moreover, for 42 problems the set of clauses produced by  $\mathcal{EC}$  does not contain constrained facts, and thus its satisfiability can immediately be checked by Z3.

Unfortunately, it may happen that our transformation prevents the solution of some verification problems. Indeed, the transformation algorithm  $\mathcal{EC}$  does not terminate within the timeout on two problems that can be proved using Z3 alone. However, on the 10 problems that are solved by both Z3 and  $\mathcal{EC}$ ;Z3, the average time taken by  $\mathcal{EC}$ ;Z3 (3.41s) to solve one problem is much lower than that taken by Z3 alone (9.74s).

Finally, let us comment on our benchmark suite. Many of the problems are small,

but the properties to be verified are not trivial, such as those presented in Sections 2 and 6. One of the most difficult problems solved by  $\mathcal{EC}$ ; Z3, but not by RCAML, consists in showing that the insertion of a node in a binary search tree produces again a binary search tree. That solution took about 5s. The solution of some of the 13 problems on which both  $\mathcal{EC}$ ; Z3 and RCAML run out of time, requires the use of suitable lemmas. For instance, one of these problems can be proved by using a lemma stating that the sum of the elements of the concatenation of two lists is equal to the sum of the elements of the first list plus the sum of the elements of the second list. The extension of our approach to support automatic lemma discovery is left for future work.

## 8 Related Work and Conclusions

We have presented a two-step approach to the verification of programs that manipulate inductively defined data structures, such as lists and trees. The first step consists in the transformation of the set of clauses that encodes the given verification problem, into a new, equisatisfiable set of clauses whose variables do no longer refer to inductive data structures. The second step consists in the application of a CHC solver to the derived set of clauses with integer and boolean constraints only. Thus, in our approach we can take full advantage of the many efficient solvers that exist for clauses with integer and boolean constraints (de Moura and Bjørner 2008, Grebenshchikov et al. 2012, Hoder and Bjørner 2012, Hojjat et al. 2012, Komuravelli et al. 2013). Through some experiments we have shown that, using an implementation of our algorithm and the Z3 solver, our two-step approach is competitive with respect to other techniques that extend CHC solvers by adding deduction rules for reasoning on inductive data structures (Unno et al. 2017).

Our transformation technique is related to methods proposed in the past for improving the efficiency of functional and logic programs, such as *deforestation* (Wadler 1990), *unnecessary variable elimination* (UVE) (Proietti and Pettorossi 1995), and *conjunctive partial deduction* (De Schreye et al. 1999). Among these techniques, the UVE transformation, which makes use of the fold/unfold rules, is the most similar to the one presented in this paper. However, in this paper we have introduced several technical novelties with respect to UVE: (i) the use of type information, (ii) the use of constraints, and (iii) a better characterization of the termination of the main algorithm (in particular, here we have introduced the notions of a slice decomposition and a circular sharing). At a more general level, the objective of the work presented in this paper is to show the usefulness of our transformation techniques for the improvement of the effectiveness of CHC solvers, rather than the improvement of the execution of logic programs.

The idea of using a transformation-based approach for the verification of software comes from the area of Constraint Logic Programming, where program specialization has been applied as a means of deriving CLP programs from interpreters of imperative languages (Albert et al. 2007, De Angelis et al. 2017b, Méndez-Lojo et al. 2008, Peralta et al. 1998). CHC solvers based on combinations of transformations and abstract interpretation have also been developed (De Angelis et al. 2014, Kafle et al. 2016) and have been shown to be competitive with solvers based on CEGAR, Interpolation, and PDR.

Recently, CHCs have been proposed for verifying *relational* program properties (Felsing et al. 2014), that is, properties that relate two programs, such as equivalence. It has also been shown that *predicate pairing*, which is a fold/unfold transformation for CHCs,

greatly improves the effectiveness of CHC solvers for relational verification (De Angelis et al. 2015, De Angelis et al. 2016). A related CHC transformation technique, called *CHC product*, works by composing pairs of clauses with an effect similar to predicate pairing, although in some cases it may derive sets of clauses with fewer predicates (Mordvinov and Fedyukovich 2017). Neither predicate pairing nor CHC product can remove inductively defined data structures, as done by the transformation technique presented in this paper.

Similarly to the technique presented in this paper, fold/unfold transformations and constraint generalization have also been used in a verification technique for imperative programs that compute over arrays (De Angelis et al. 2017a). However, the above mentioned technique is not able to remove array data structures, and unlike the one presented here, it does not consider inductively defined data structures.

We plan to extend our transformation-based verification method in a few directions, and in particular we plan: (i) to study the problem of automatically generating the lemmas which are sometimes needed for removing data structures, and (ii) to consider the verification problem for higher-order functional programs.

# 9 Acknowledgements

We would like to thank Hiroshi Unno for his support in the use of the RCAML system. We thank the anonymous reviewers for their constructive comments. The authors are members of the INdAM Research group GNCS. E. De Angelis, F. Fioravanti and A. Pettorossi are research associates at CNR-IASI, Rome, Italy.

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## Appendix

In order to prove Theorem 2, we first recall some notions and results regarding the transformation rules and their correctness.

A transformation sequence is a sequence  $S_0, S_1, \ldots, S_n$  of sets of CHCs, whose constraints are in  $LIA \cup Bool$ , where, for  $i=0,\ldots,n-1$ ,  $S_{i+1}$  is derived from  $S_i$  by applying one of the following rules R1–R5.

Let  $Defs_i$  denote the set of all the clauses, called *definitions*, introduced by rule R1 during the construction of the transformation sequence  $S_0, S_1, \ldots, S_i$ . In particular,  $Defs_0 = \emptyset$ .

(R1) Define. We introduce a clause  $D: newp(X_1, \ldots, X_k) \leftarrow c, G$ , where: (i) newp is a predicate symbol not occurring in the sequence  $S_0, S_1, \ldots, S_i$ , (ii) c is a constraint in  $LIA \cup Bool$ , (iii) G is a non-empty conjunction of atoms whose predicate symbols occur in  $S_0$ , and (iv)  $X_1, \ldots, X_k$  are distinct variables occurring in (c, G). Then, we derive the new set  $S_{i+1} = S_i \cup \{D\}$  and  $Defs_{i+1} = Defs_i \cup \{D\}$ .

(R2) Unfold. Let  $C: H \leftarrow c, L, A, R$  be a variant of a clause in  $S_i$ . Let  $K_1 \leftarrow c_1, B_1, \ldots, K_m \leftarrow c_m, B_m$  be all clauses of  $S_i$  (without loss of generality, we assume  $vars(S_i) \cap vars(C) = \emptyset$ ) such that, for  $j = 1, \ldots, m$ , (1) there exists a most general unifier  $\vartheta_j$  of A and  $K_j$ , and (2) the constraint  $(c, c_j)\vartheta_j$  is satisfiable. By unfolding the atom A in C using  $S_i$  we derive the new set  $S_{i+1} = (S_i \setminus \{C\}) \cup \{(H \leftarrow c, c_1, L, B_1, R)\vartheta_1, \ldots, (H \leftarrow c, c_m, L, B_m, R)\vartheta_m\}$ .

(R3) Fold. Let  $C: H \leftarrow c, L, Q, R$  be a clause in  $S_i$ , where Q is a non-empty conjunction of atoms, and let  $D: K \leftarrow d, B$  be (a variant of) a clause in  $Defs_i$  with  $vars(C) \cap vars(D) = \emptyset$ . Suppose that there exist a substitution  $\vartheta$  and a constraint e such that: (i)  $Q = B\vartheta$ , (ii)  $LIA \cup Bool \models \forall (c \leftrightarrow (e \land d\vartheta))$ , and (iii) for every variable  $X \in vars(d, B) \setminus vars(K)$ , the following conditions hold: (iii.1)  $X\vartheta$  is a variable not occurring in  $\{H, c, L, R\}$ , and (iii.2)  $X\vartheta$  does not occur in the term  $Y\vartheta$ , for any variable Y occurring in (d, B) and different from X. By folding C using the definition D, we derive clause  $E: H \leftarrow e, L, K\vartheta, R$ . In this case we also say that E is derived by folding Q in C. We derive the new set  $S_{i+1} = (S_i \setminus \{C\}) \cup \{E\}$ .

(R4) Replace Equivalent Constraints. Let us consider a subset of  $S_i$  of the form  $\{(H \leftarrow c_1, G), \dots, (H \leftarrow c_k, G)\}$ . Suppose that, for some constraints  $d_1, \dots, d_m$ ,

$$LIA \cup Bool \models \forall (\exists Y_1 \dots \exists Y_r \ (c_1 \lor \dots \lor c_k) \leftrightarrow \exists Z_1 \dots \exists Z_s \ (d_1 \lor \dots \lor d_m))$$

where  $\{Y_1, \ldots, Y_r\} = vars(c_1 \lor \ldots \lor c_k) \setminus vars(\{H,G\})$  and  $\{Z_1, \ldots, Z_s\} = vars(d_1 \lor \ldots \lor d_m) \setminus vars(\{H,G\})$ . Then, we derive the new set  $S_{i+1} = (S_i \setminus \{(H \leftarrow c_1, G), \ldots, (H \leftarrow c_k, G)\}) \cup \{(H \leftarrow d_1, G), \ldots, (H \leftarrow d_m, G)\}.$ 

Note that rule R4 enables the deletion of a clause with an inconsistent constraint in its body. Indeed, if  $c_1$  is unsatisfiable, then  $LIA \cup Bool \models \forall (c_1 \leftrightarrow d_1 \lor \ldots \lor d_m)$  with m = 0.

(R5) Replace Functional Predicates. Let  $C: H \leftarrow c, G_1, p(t, u), G_2, p(t, w), G_3$ , be a clause in  $S_i$  and let p(X, Y) be functional in  $S_i$  (see Definition 3). Then, we derive the new set  $S_{i+1} = (S_i \setminus \{C\}) \cup \{(H \leftarrow c, G_1, p(t, u), G_2, G_3)\vartheta\}$ , where  $\vartheta$  is the most general unifier of u and w.

The following theorem sums up various results presented in the literature (Etalle and Gabbrielli 1996, Tamaki and Sato 1984).

## Theorem 4 (Equivalence with respect to satisfiability)

Let  $S_0, S_1, \ldots, S_n$  be a transformation sequence such that every definition in  $Defs_n$  is unfolded during the construction of this sequence. Then,  $S_0 \cup LIA \cup Bool$  is satisfiable if and only if  $S_n \cup LIA \cup Bool$  is satisfiable.

Now, we prove Theorems 2 and 3 of Section 5.

## Theorem 2 (Partial Correctness)

Let Cls be a set of definite clauses and let Gs be a set of goals. If Algorithm  $\mathcal{E}$  terminates for the input clauses  $Cls \cup Gs$ , returning a set TransfCls of clauses, then (1)  $Cls \cup Gs$  is satisfiable iff TransfCls is satisfiable, and (2) all clauses in TransfCls have basic types. *Proof* 

Point (1) follows from Theorem 4 by taking into account that: (i) Algorithm  $\mathcal{E}$  can be viewed as a particular sequence of applications of Rules R1–R5, and (ii) every definition in *Defs* is unfolded during the execution of  $\mathcal{E}$ .

Point (2) follows from the fact that, by construction, every clause introduced in *TransfCls* has basic types. To see this, let us consider a clause C in *TransfCls*. Clause C belongs to the set *FldCls* of clauses obtained by a *Define-Fold* step. Looking at the *Define-Fold* procedure, we have that: (i) the head of C is either *false* (because  $C \in Gs$ ) or its head predicate has been introduced by the *Define* step, and hence, by construction, has basic types, and (ii) the body of C has the form:  $c, newp_1(V_1), \ldots, newp_n(V_n)$ , where c is a constraint which has basic types (because it belongs to  $LIA \cup Bool$ ) and  $newp_1, \ldots, newp_n$  are predicates that, by construction, have basic types.

## Theorem 3 (Termination)

Let Cls be a set of definite clauses such that every clause in Cls has a disjoint, quasidescending slice decomposition. Let Gs be a set of goals such that, for each goal  $G \in Gs$ , (i) G is of the form  $false \leftarrow c, A_1, \ldots, A_m$ , where for  $i = 1, \ldots, m$ ,  $A_i$  is an atom whose arguments are distinct variables, and (ii) G has no sharing cycles. Then Algorithm  $\mathcal{E}$ terminates for the input clauses  $Cls \cup Gs$ .

## Proof (Sketch)

Without loss of generality, we assume that for every clause  $H \leftarrow c, B$  every variable of basic type has at most one occurrence in H, B.

For any tree t, by height(t) we denote the height of t, that is, the maximal length of a path from the root of t to one of its leaves. We extend the function height to terms and atoms, viewed as trees. First of all, we observe that Algorithm  $\mathcal{E}$  terminates iff there exists two non-negative integers H and N such that, for every definition  $newp(V) \leftarrow A_1, \ldots, A_n$  added to *Defs* during the execution of  $\mathcal{E}$ , we have that, for  $i = 1, \ldots n$ ,  $height(A_i) \leq H$ , and  $n \leq N$ . Indeed, the existence of H and N implies the finiteness of the set of definitions introduced by  $\mathcal{E}$ , and hence the finiteness of the number of iterations of the body of the while-do loop of  $\mathcal{E}$  itself.

Let us consider a clause  $D: newp(V) \leftarrow A_1, \ldots, A_n$  added to *Defs* during the execution of  $\mathcal{E}$ . Then D satisfies the following properties:

- P1. The goal  $false \leftarrow A_1, \ldots, A_n$  obtained from D by replacing newp(V) by false, has no sharing cycles;
- P2. All atoms in the body of D are linear;
- P3. For any two distinct atoms  $A_i$  and  $A_j$  in the body of D, if  $A_i$  and  $A_j$  share a

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non-basic variable, then they are of the form  $p(\ldots,t_i,\ldots)$  and  $q(\ldots,t_j,\ldots)$ , where either  $t_i \leq t_j$  or  $t_j \leq t_i$ , and  $A_i$  and  $A_j$  do not share any variable besides the ones in  $vars(t_i) \cap vars(t_j)$ ;

- P4. For any atom  $A_i$  in the body of D,  $height(A_i) \leq H$ , where H is the maximal height of an atom in  $Cls \cup Gs$ ;
- P5. Let  $V_G$  be the number of occurrences of non-basic variables in a goal G, and let M be  $\max\{V_G \mid G \in Gs\}+1$ . Then, in the body of D, (P5.1) every non-basic variable has at most M occurrences, and (P5.2) there exist  $K \leq M$  predicate arguments such that every non-basic variable that occurs more than once, also occurs in one of those K arguments.

Property P1 holds for each clause D initially in *Defs* by the hypothesis that Gs is a set of goals that have no sharing cycles. This property, when referred to the body of the clauses, is preserved by the *Unfold* and *Replace* steps, due to the hypothesis on the clauses in *Cls*, and hence it also holds for each new definition added to *Defs* by the *Define* step after *Unfold* and *Replace*.

Property P2 holds for each clause D initially in *Defs* by Hypothesis (i) on *Gs*. This property is preserved by the *Unfold* and *Replace* steps, due to the hypothesis on the clauses in *Cls*. Note, in particular, that the existence of a disjoint, quasi-descending slice decomposition for all clauses in *Cls* implies that each atom in the body of a clause in *Cls* is linear, and hence only linear atoms are introduced by *Unfold* steps. The linearity of the atoms different from the one replaced by an *Unfold* step is enforced by the existence of a disjoint, quasi-descending slice decomposition for all clauses in *Cls*. Linearity is also preserved by *Replace* steps. Thus, Property P2 follows from the fact that the body of Dconsists of atoms taken from the body of a clause derived by *Unfold* and *Replace* steps.

Property P3 holds for each clause D initially in *Defs* by Hypothesis (i) on *Gs*. Property P3 also holds for each clause derived by the *Unfold* and *Replace* steps by Property P1 and by the hypothesis that every clause in *Cls* has a disjoint, quasi-descending slice decomposition. Then, Property P3 follows from the fact that the body of D consists of atoms taken from the body of clauses derived by *Unfold* and *Replace* steps.

Property P4 holds for each clause D initially in *Defs* because the body of clause D is the set of atoms occurring in the body of a goal in Gs. This property also holds for each clause derived by Unfold steps. Indeed, suppose that we unfold an atom A in the clause C of the form  $H \leftarrow c, L, A, R$  such that either (i) A is strictly maximal in L, A, R, or (ii) all atoms in L, A, R are not strictly maximal. Both in case (i) and case (ii), by Property P3, A is of the form  $p(\ldots, t_i, \ldots)$  and any atom Q in L,R that shares a non-basic variable with A is of the form  $q(\ldots, t_j, \ldots)$ , with  $t_j \leq t_i$ , and A and Q do not share any variable besides the ones in  $vars(t_i) \cap vars(t_j)$ . Let  $K_1 \leftarrow c_1, B_1, \ldots, K_m \leftarrow c_m, B_m$  be all clauses of Cls (with  $vars(Cls) \cap vars(C) = \emptyset$ ) such that, for  $i = 1, \dots, m$ , (i) there exists a most general unifier  $\vartheta_i$  of A and  $K_i$ , and (ii) the constraint  $(c, c_i)\vartheta_i$  is satisfiable. Then, by unfolding A in C we derive the clauses  $(H \leftarrow c, c_1, L, B_1, R)\vartheta_1, \dots, (H \leftarrow c, c_m, L, B_m, R)\vartheta_m$ . By Property P2 A is a linear atom, and by the hypothesis that every clause in  $Cl_s$  has a disjoint, quasi-descending slice decomposition, we have that, for i = 1, ..., m, for every atom E in L,  $B_i, R$ ,  $height(E\vartheta_i) \leq \max(\{height(E), height(A), height(K_i)\})$ . Thus, if Property P4 holds for C, then it also holds for the clauses  $(H \leftarrow c, c_1, L, B_1, R)\vartheta_1, \ldots,$  $(H \leftarrow c, c_m, L, B_m, R)\vartheta_m$ . Property P4 also holds for each clause derived by *Replace* steps,

and hence it also holds for D, whose body consists of atoms taken from the body of clauses derived by *Unfold* and *Replace* steps.

Property P5 holds for each clause D initially in *Defs* because the body of clause D is the set of atoms occurring in the body of a goal G in Gs for which Hypothesis (i) holds. Now we prove that the following two properties, which generalize Property P5, hold for each clause E derived by an *Unfold* step: in each sharing block in the body of E, (P5.1) every non-basic variable has at most M occurrences, and (P5.2) there exist  $K \leq M$  predicate arguments such that every non-basic variable with more than one occurrence in the body of E, also occurs in one of those K arguments.

Suppose that P5.1 and P5.2 hold for a clause C of the form  $H \leftarrow c, L, A, R$ , and we unfold A in C. Let  $K_1 \leftarrow c_1, B_1, \ldots, K_m \leftarrow c_m, B_m$  be all clauses of Cls (with  $vars(Cls) \cap vars(C) = \emptyset$ ) such that, for  $i=1,\ldots,m$ , (i) there exists a most general unifier  $\vartheta_i$  of A and  $K_i$ , and (ii) the constraint  $(c,c_i)\vartheta_i$  is satisfiable. Then, by unfolding A in C we derive the clauses  $C_1: (H \leftarrow c, c_1, L, B_1, R)\vartheta_1, \ldots, C_m: (H \leftarrow c, c_m, L, B_m, R)\vartheta_m$ . By Properties P2 and P3, and by the existence of a disjoint, quasi-descending slice decomposition, for  $i=1,\ldots,m$ , the number of occurrences of any variable with more than one occurrence in the body of  $C_i$ , is not larger than the maximal number of occurrences of any variable in the body of  $C_i$ , and hence Property P5.1 holds for the body of  $C_i$ .

Moreover, suppose that in every sharing block in the body of C there exist  $K \leq M$ arguments  $t_1, \ldots, t_K$  such that every variable variable with more than one occurrence, also occurs in one of those K arguments. By Property P3, we may assume that  $t_1, \ldots, t_K$ are maximal with respect to the  $\leq$  relation and do not share any variable. Looking at the Unfold procedure, the atom A selected for unfolding must have one among  $t_1, \ldots, t_K$ as an argument, say  $t_1$ . By our hypotheses, if by unfolding A the argument  $t_1$  is replaced by more than one term, these new terms must appear in different sharing blocks, and hence the number of maximal arguments in each sharing block does not increase. Thus, Property P5.2 holds for  $C_1, \ldots, C_m$ . Similarly, we can prove that Properties P5.1 and P5.2 hold for each clause derived by *Replace* steps, and hence Property 5 holds for D, whose body consists of a sharing block of a clause derived by Unfold and Replace steps.

Now, from Properties P4 and P5 it follows that there exists an integer J, depending on H and M, such that in the body of D there are at most J distinct variables. Thus, there exists N such that the body of D has at most N atoms of height not larger than H, and hence the thesis holds.