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#### Abstract

In this paper we describe how induction formulas are formulated in verification of Prolog programs. The same problem for well-founded induction was investigated by Boyer and Moore in their verification system for terminating LISP programs (BMTP). The least fixpoint semantics of Prolog provides an advantage that computational induction is easily applicable and generates simpler induction schemes. We investigate how the computational induction is applied to a class of first order formulas called S-formulas, in which specifications are described in our verification system. In addition, we show how an equivalence preserving transformation can be utilized to merge induction schemes into simpler one when more than two induction schemes are suggested.

Keywords: Program Verification, Induction, Prolog, Program Transformation.

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#### 1. Introduction

The intimacy of Prolog to first order logic is expected to bring advantages to first order inference in verification ([11],[17]). But how about induction, which plays an important role in verification of programs? The least fixpoint semantics of Prolog suggests fixpoint inductions. But how can we find such induction schemes from theorems to be proved? In addition, when more than two different induction schemes are suggested, how can we manage them?

In this paper we answer these questions. So far the same problem was investigated by Boyer and Moore [3] in their verification system for terminating LISP programs (BMTP) where they adopted quantifier-free formulas as specifications and applied well-founded induction. The least fixpoint semantics of Prolog provides an advantage that computational induction is applicable staying within first order logic and moreover it generates simpler induction schemes [5]. Here we investigate how the computational induction is applied to a class of first order formulas called S-formulas, in which specifications are described in our verification system. In addition, we show how the Tamaki-Sato's Prolog program transformation can be utilized to merge computational induction schemes into more simpler one when more then two induction schemes are suggested. Because the Tamaki-Sato's transformation preserves equivelence, justification of the manipulation of induction schemes is immediate and easy to grasp the meaning.

After summarizing preliminary materials in section 2 and a framework of verification in section 3, we describe in section 4 how computational induction schemes are found from theorems to be proved. In section 5, we show how induction schemes are merged into a more appropriate one when more than two induction schemes are suggested. Lastly in section 6 we discuss the relations to other works and our actual verification system.

#### 2. Preliminaries

#### 2.1. Polarity of Subformulas

We generalize the distinctions of positive and negative subformulas. The positive and negative subformula of a formula  $\mathcal{F}$  is defined as follows (Murray [14]).

- (a) \$\mathcal{I}\$ is a positive subformula of \$\mathcal{I}\$.
- (b) When ¬G is a positive (negative) subformula of F, then G is a negative (positive) subformula of F.
- (c) When G∧H or G∨H is a positive (negative) subformula of F, then G and H are positive (negative) subformulas of F.

- (d) When  $\mathcal{G} \supset \mathcal{H}$  is a positive (negative) subformula of  $\mathcal{F}$ , then  $\mathcal{G}$  is a negative (positive) subformula of  $\mathcal F$  and  $\mathcal H$  is a positive (negative) subformula of  $\mathcal F$ .
- (e) When  $\forall X \mathcal{G}, \exists X \mathcal{G}$  are positive (negative) subformulas of  $\mathcal{F}$ , then  $\mathcal{G}_X(t)$  is a positive (negative) subformula of  $\mathcal{F}$ .

Example 2.1: Let 7 be

∀ A,B (reverse(A,B) ⊃reverse(B,A)).

Then reverse(A, B) is a negative subformula of  $\mathcal{F}$ .

## 2.2. S-formulas and Goal Formulas

Let  $\mathcal F$  be a closed first order formula. When  $\forall X \mathcal G$  is a positive subformula or  $\exists X \mathcal G$ is a negative subformula of  $\mathcal{F}$ , X is called a free variable of  $\mathcal{F}$ . When  $\forall Y \mathcal{H}$  is a negative subformula or  $\exists Y \ \forall$  is a positive subformula of  $\mathcal{F}$ , Y is called an undecided variable of  $\mathcal{F}$ . In other words, free variables are variables quantified universally and undecided variables are those quantified existentially when  $\mathcal F$  is converted to prenex normal form.

Example 2.2.1: Let F be

 $\forall A,V,C (append(\Lambda,[V],C) \supset \exists B reverse(C,[V|B])).$ 

Then A, V, C are all free variables, while B is an undecided variable.

A closed first order formula S is called an S-formula when

- (a) no free variable in S is quantified in the scope of quantification of an undecided variable in S and
- (b) no undecided variable appears in negative atoms of S.

In other words, S-formulas are formulas convertible to prenex normal form  $\forall X_1, X_2, \ldots, X_n$  $\exists Y_1, Y_2, \ldots, Y_m \mathcal{F}$  and no  $Y_1, Y_2, \ldots, Y_m$  appears in negative atoms of  $\mathcal{F}$ . Note that S-formulas include both universal formulas  $\forall X_1, X_2, \dots, X_n \mathcal{F}$  and usual execution goals  $\exists Y_1, Y_2, \dots, Y_m$  $(A_1 \wedge A_2 \wedge \cdots \wedge A_k).$ 

Example 2.2.2: Let S be

 $\forall A,V,C (append(A,[V],C) \supset \exists B reverse(C,[V|B])).$ 

Then S is an S-formula, because free variables A, V, C are quantified outside  $\exists B$  and B appears only in the positive atom reverse(C, [V|B]). A universal formula  $\forall A, B$  (reverse(A, B)  $\supset$ reverse(B,A)) and an execution goal  $\exists Cappend([1,2],[3],C)$  are also S-formulas.

A formula G obtained from an S-formula S by replacing free variable X with X, undecided variable Y with ?Y and deleting all quantifications is called a goal formula of S. Note that S can be uniquely restorable from G. In the followings, we use goal formulas in stead of original S-formulas with explicit quantifiers. Goal formulas are denoted by F, G, H.

Example 2.2.3: An S-formula

 $\forall$  A,V,C (append(A,[V],C)  $\supset \exists$ B reverse(C,[V|B])).

is represented by a goal formula

 $append(\Lambda,[V],C)\supset reverse(C,[V|?B]).$ 

A universal formula  $\forall A, B(reverse(A,B) \supset reverse(B,A))$  and an execution goal  $\exists Cappend([1,2],[3],C)$ are represented by  $reverse(A, B) \supset reverse(B, A)$  and append([1, 2], [3], ?C) respectively.

# 2.3. Manipulation of Goal Formulas

Lastly we introduce two manipulations of goal formulas. One is an application of a class

of substitutions. To avoid variable name's conflict, we introduce a class of substitutions. A substitution  $\sigma$  is called a *substitution away from A* when  $\sigma$  instanciates each free variable X in A to t such that every variable in t is a fresh free variable not in A.

Example 2.3.1:  $\langle A \leftarrow [X|L], B \leftarrow M \rangle$  is a substitution away from reverse(A, B) and unifies it with reverse([X|L], M), where X, L and M are considered fresh free variables.  $\langle C \leftarrow [X|L], U \leftarrow Y, B \leftarrow [X|N] \rangle$  is a substitution away from append(C, [U], B) and unifies it with append([X|L], [Y], [X|N]) where X, L, Y and N are considered fresh free variables.

Another manipulation is a reduction of goal formulas with logical constants true and false. The reduced form of a goal formula G, denoted by  $G \downarrow$ , is the normal form in the reduction system defined as follows.

Example 2.3.2: Let  $G_1$  and  $G_2$  be false  $\supset$  reverse (B, A) and true  $\supset$  reverse (B, A). Then  $G_1 \downarrow$  is true and  $G_2 \downarrow$  is reverse (B, A).

#### 3. Framework of Verification of Prolog Programs

#### 3.1. Programming Language

We introduce type construct into Prolog to separate definite clauses defining data structures from others defining procedures, e.g.,

```
type.
    list([]).
    list([X|L]) := list(L).
end.
```

The body of type is a conjunction of definite clauses whose head is with a unary predicate defining a data structure. (Type predicates in our verification system are being corresponded to shells in BMTP.) Procedures are defined by following the syntax of DEC-10 Prolog [15],e.g.,

```
\begin{array}{l} \operatorname{append}([\ ],K,K), \\ \operatorname{append}([X|L],M,[X|N]) := \operatorname{append}(L,M,N), \\ \operatorname{reverse}([\ ],[\ ]), \\ \operatorname{reverse}([X|L],M) := \operatorname{reverse}(L,N), \operatorname{append}(N,[X],M). \end{array}
```

Throughout this paper, we study pure Prolog consisting of definite clauses and consider a finite set of definite clauses P as their conjunction. We assume variables in each definite clause are renamed at each use so that there occurs no variable names conflict.

#### 3.2. Specification Language

The main construct of our specification language is theorem to state a theorem to be proved, e.g.,

theorem(halting-theorem-for-append). ∀ A:list,B ∃ C append(A,B,C). end.

The body of theorem must be a closed S-formula. Any variable X in quantification may be followed by a type qualifier : p (e.g.,:list above).  $\forall X:p\mathcal{F}$  and  $\exists X:p\mathcal{F}$  are abreviations of  $\forall X(p(X)\supset \mathcal{F})$  and  $\exists X(\mathcal{F}\land p(X))$  respectively.

## 3.3. Framework of Verification

Let S be a specification in an S-formula,  $M_0$  be the minimum Herbrand model of Pand  $P^*$  be the completion of P. We adopt a formulation that verification of S with respect to P is to show  $M_0 \models S$  when model theoretically speaking and to prove S from P using the first order inference and some induction when proof theoretically speaking.

The most important difference between our system and BMTP is that specifications in BMTP are quantifier-free (i.e. universal) formulas while ours are general first order formulas. Though we prove quantifier-free specifications of the form  $\forall X_1, X_2, \ldots, X_n$   $(A_1 \land A_2)$  $\wedge \cdots \wedge A_m \supset A_0$ ) in most cases, the consideration of existential quantifiers is inevitable because of the effects of internal variables in Prolog. For example, suppose we prove  $\forall X, Y (condition(X, Y))$  $\supset p(X,Y)$ ) with respect to a program p(X,Y) = q(X,Z), r(Z,Y). Then we must prove  $\forall X,Y$  $(condition(X,Y) \supset \exists Z(q(X,Z) \land r(Z,Y))$  substantially.

# 4. Generation of Computational Induction Schemes

## 4.1. Computational Induction

For pure Prolog, the transformation T of Herbrand interpretations is always continuous and there holds  $\bigcup_{i=0}^{\infty} T^i(0) = M_0$ . This suggests use of fixpoint induction to prove  $M_0 \models \psi$ . Here we explain it rather intuitively following Clark [5] p.75-76.

Example 4.1.1: Let reverse be a relation defined by the previous definite clauses. The reverse relation is the smallest set of pairs of terms that includes a pair ([], []) and that, for any term s,includes  $([s|t_1],t_2)$  whenever it includes  $(t_1,t)$  and  $(t,[s],t_2)$  is in the append relation. Hence, suppose Q(A,B) is a formula with free variable A,B. For any Herbrand interpretation, Q(A,B) will denote some binary relation over terms. If this relation includes ([],[]),i.e.

is true, and if it includes  $([s|t_1], t_2)$  whenever it includes  $(t_1, t)$  and  $(t, [s], t_2)$  is in append relation, i.e.,

 $\forall \ A,B,C,U \ (Q(A,C) \land append(C,[U],B) \supset Q([U|A],B))$ is true, then the relation Q(A,B) includes all the pairs of terms in the reverse relation. In

 $\forall A,B (reverse(A,B) \supset Q(A,B))$ is true of the reverse relation and such Q(A,B). Hence we get the following computational induction scheme

$$\frac{Q([\ ],[\ ])}{\forall A,B,C,U} \frac{(Q(A,C)\land append(C,[U],B)) \supset Q([U|A],B))}{\forall A,B}$$

$$\frac{\forall A,B \text{ (reverse(A,B))} \supseteq Q(A,B))}{\forall A,B}$$

This is a Prolog version of the de Bakker and Scott's computational induction [2]. In order to apply such an induction, we need to know (a) how we select the key atom (reverse(A, B) above), (b) what conditions we have to observe for soundness and (c) how we can make the generated induction schemes simple.

First of all, we introduce a concept "dominant atom". An atom A in a goal formula G is said to be dominant in G when G is equivalent to a goal formula of the form  $A \supset H$ . Note that A is dominant in G iff A is negative and  $G_A[false] \downarrow$  is true. Then H is  $G_A[true] \downarrow$ .

Example 4.1.2: Let G be  $reverse(A, B) \supset reverse(B, A)$ . Then the left reverse(A, B) is dominant in G, but the right reverse(B, A) is not.

#### 4.2. Inducible Definite Clause

A definite clause is said to be inducible to A when, for any ground instance of the definite clause such that the head is a ground instance of A, any recursive call in the body is also a ground instance of A. (Hence any non-recursive definite clause is always inducible. Any definite clause with a head nonunifiable with A is also inducible.)

When every definite clause in P is inducible to A, A is said to be closed with respect to P. This means that the set of ground atoms in  $M_0$  of the form of instance of A is computable by the instances of inducible definite clauses. Note that an atom  $p(X_1, X_2, ..., X_n)$  in general form is always closed.

Example 4.2.1: Let the atom A be append(A,  $\{U\}$ , C). Then A is closed. This means that  $\{append(t_1, [t_2], t_3) \mid t_1, t_2 \text{ and } t_3 \text{ are ground terms}\} \cap M_0$  is computable by some instances of definite clauses, i.e.,

```
append([],[Y],[Y]).

append([X|L],[Y],[X|N]) := append(L,[Y],N).
```

The inducibility can be checked as follows.

- (a) Check whether the head B<sub>0</sub> is unifiable with A by a substitution for A away from A (see 2.3). If it is, decompose the mgu to σ ∘ τ<sub>0</sub> where σ is the restriction to variables in B<sub>0</sub> and τ<sub>0</sub> is the restriction to variables in A. If it is not, the definite clause is inducible to A.
- (b) Check whether each instance of the recursive call in the body σ(B<sub>i</sub>) is an instance of A and if it is, compute the instanciation τ<sub>i</sub>. If it is not, the definite clause is not inducible to A.

The set of all instances of definite clauses by  $\sigma$  is called instanciated program for A.

Example 4.2.2: Let the atom A be append(A, [U], C). Then the first head append([], L, L) is unifiable with append(A, [U], C) by  $< L \Leftarrow [Y] > \circ < A \Leftarrow [], U \Leftarrow Y, C \Leftarrow [Y] >$ . The second head append([X|L], M, [X|N]) is unifiable with append(A, [U], C) by  $< M \Leftarrow [Y] > \circ < A \Leftarrow [X|L], U \Leftarrow Y, C \Leftarrow [X|N]$  and the instance append(L, [Y], N) in the body is also an instance of append(A, [U], C) by  $< A \Leftarrow L, U \Leftarrow Y, C \Leftarrow N >$ .

#### 4.3. Generation of Induction Schemes

Let G be a goal formula,  $A=p(t_1,t_2,...,t_n)$  be a dominant atom in G and Q be  $G_A[true] \downarrow$ . Let "B<sub>0</sub> :- B<sub>1</sub>, B<sub>2</sub>,..., B<sub>m</sub>" be a definite clause inducible to A. By  $\sigma(B_0 :- B_1, B_2,..., B_m)$ " be a definite clause inducible to A. By  $\sigma(B_0 :- B_1, B_2,..., B_m)$ " be a definite clause inducible to A.

 $B_1, B_2, \ldots, B_m)_p(Q)$  we denote a formula obtained by replacing  $\sigma(B_i) = p(s_1, s_2, \ldots, s_n)$  in  $\sigma(B_1 \wedge B_2 \wedge \cdots \wedge B_m \supset B_0)$  with  $\tau_i(Q)$  when  $B_0$  is unifiable with A and true otherwise.

We generate an induction scheme as follows. All free variables are quantified universally at the outermost, which keeps the generated subgoals within S-formulas.

at the outermost, which keeps the generators 
$$G$$
  $\sigma_1(B_{10}:-B_{11},B_{12},\ldots,B_{1m_1})_p(Q), \ \sigma_2(B_{20}:-B_{21},B_{22},\ldots,B_{2m_2})_p(Q), \ \ldots, \ \sigma_k(B_{k0}:-B_{k1},B_{k2},\ldots,B_{km_k})_p(Q)$ 

Example 4.3.1: Suppose we prove a theorem

theorem(reverse-reverse).

 $\forall A,B \; (\texttt{reverse}(A,B) \supset \texttt{reverse}(B,A)).$ 

Then reverse(A, B) is dominant. Let Q(A, B) be  $(true \supset reverse(B, A)) \downarrow = reverse(B, A)$ . The induction scheme generated is the one in example 4.1,i.e.,

reverse([],[]) 
$$\forall A,B,C,U \text{ (reverse(C,A) } \land append(C,[U],B) } \supset reverse(B,[U|A]))$$

$$\forall A,B \text{ (reverse(A,B) } \supset reverse(B,A))$$

### 4.4. Generalization

Must we always give up applications of computational induction when some definite clause is not inducible?

Example 4.4.1: Let A be reverse(A, [V|B]). Then "reverse([X|L], M) :- reverse(L, N), append(N, [X], M)" is not inducible to A, because the instance of the recursive call by  $\sigma$  in the definition above may have the form reverse(L, N) which is not necessarily an instance of reverse(A, [V|B]).

A set of replacement M of occurrences of subterm in A with fresh free variables is called a generalization mask when no different subterm is replaced with a same fresh free variable. An atom obtained from A by replacing the occurrence of  $t_{i_k}$  with a fresh free variable  $X_{i_k}$  for all  $< t_{i_k} = X_{i_k} > \in M$  is called generalization of A with respect to M and denoted by  $A_M$ . Then  $X_{i_1} = t_{i_1} \wedge X_{i_2} = t_{i_2} \wedge \cdots \wedge X_{i_k} = t_{i_m}$  is called the generalization equation. (When  $M = \emptyset$ ,  $A_{\emptyset}$  is A itself and the generalization equation is true.)

Example 4.4.2:  $M = \{ [V|B] \leftarrow B' \}$  is a generalization mask of reverse(A, [V|B]) in  $\forall$  A,V,B (reverse(A,[V|B])  $\supset$  member(V,A)).

The generalization of reverse(A, [V|B]) w.r.t. M is reverse(A, B') where B' is a fresh free variable. The generalization equation is B' = [V|B]. Intuitively this corresponds to modify the theorem to

 $\forall A,V,B,B' \text{ (reverse}(A,B') \land B' = [V|B] \supset member(V,A)).$ 

A definite clause  $B_0: B_1, B_2, ..., B_m$  is said to be inducible with M to A when it is inducible to  $A_M$ .

Example 4.4.3: Let A be reverse(A, [V|B]) and M be  $\{[V|B] \Leftarrow B'\}$ . Then "reverse([X|L], M) :- reverse(L, N), append(N, [X], M)" is inducible with M to A, because the head reverse([X|L], M) is unifiable with reverse(A, B') by  $<> \circ < A \Leftarrow [X|L], B' \Leftarrow M >$  and the recursive call reverse(L, N) in the body is also an instance of reverse(A, B').

One naturally expect that  $A_{M}$  be the most specific one. An atom  $\overline{A}$  is said to be a

closure of A with respect to P when

- (a) A is closed with respect to P,
- (b) A is an instance of A and
- (c)  $\overline{A}$  is an instance of any  $\overline{A}'$  satisfying (a) and (b).

The closure is unique up to renaming and A is closed iff  $A = \overline{A}$  modulo renaming. (See appendix for the proof of uniqueness and algorithm to compute the closure.) A generalization mask M corresponding to  $\overline{A}$  is called a *computational induction mask*. Examples in 4.1—3 are cases with the mask  $\emptyset$ .

Example 4.4.4: Let A be reverse(A, [V|B]) and M be  $\{[V|B] \leftarrow B'\}$ . Then the first head reverse( $[\cdot], [\cdot]$ ) is unifiable with reverse(A, B') by  $<> \circ < A \leftarrow [\cdot], B' \leftarrow [\cdot] >$ . The second definite clause is inducible with M to A. Moreover reverse(A, B') is a closure of reverse(A, [V|B]) and M is a computational induction mask.

Then computational induction schemes are generated as are in 4.3 by replacing A with  $A_M$  and Q with  $G_A[\epsilon] \downarrow$  where  $\epsilon$  is the generalization equation.

Note the dominant atom reverse(A, [V|B]). Because  $\emptyset$  is not a computational induction mask, we can't apply the induction immediately without generalization. We need a non-empty mask  $\mathcal{M}=\{\ [V|B] \leftarrow B'\ \}$  and generated formulas are

```
\forall V.B ([] = [V|B] \supset member(V,[])),
```

 $\forall U, V, A, B, C, D ((C=[V|B] \supset member(V,A)) \land append(C,[U],D) \supset (D=[V|B] \supset member(V,[U|A]))).$  Note that the second subgoal remains within S-formulas. The first goal is reduced to true by the definition Z=Z. The second goal is reduced to

 $\forall U,V,A,B,C ((C=[V|B] \supset member(V,A)) \land append(C,[U],[V|B]) \supset member(V,[U|A])).$  and it is proved using the definition of append.

#### 4.5. Examples of Induction Schemes

We show how computational inductions are applied by several examples.

Example 4.5.1: We need not to impose the termination condition, i.e.  $p(t_1, t_2, ..., t_n)$  may be neither true nor false for some ground terms  $t_1, t_2, ..., t_n$ . This means we can separate  $M_0$  from other models of  $P^*$  and show properties specific to  $M_0$ . Let p and q be defined by the following programs (Apt and van Emden [1]).

```
p(a) := p(X), q(X), p(s(X)) := p(X), q(b), q(s(X)) := q(X).
```

The execution of  $? - p(s^i(a))$  never terminates, while those of  $? - p(s^j(b))$  and  $? - q(s^i(a))$  fail finitely. The minimum (and the maximum) Herbrand model is  $M_0 = \{q(b), q(s(b)), q(s(s(b))), \ldots\}$ . But we can't infer  $-p(s^i(a))$  from the completion  $P^*$ . This is showen by existence of a non-

Herbrand model M. Let the domain of M be a set consisting of red nutural numbers, green natural numbers and blue integers and the interpretation of symbols on M be one interpreting a as red 0, b as green 0, s be a function mapping X to X+1 in the usual arithmetic, p(i) is true iff i is either red natural number or blue integer and q(j) is true iff j is either green natural number or blue integer. Then M is a model of  $P^*$  not isomorphic to  $M_0$  and  $p(s^i(a))$  is valid in M (cf.Jaffar et al [10]). Suppose we prove a theorem

theorem(p-does-not-hold).

 $\forall A \neg p(A)$ .

which is valid in  $M_0$  but not valid in M, hence not provable by first order inference from  $P^*$ . Note the dominant atom p(A). Because each definite clause is inducible with  $\emptyset$ , we can apply the induction immediately and two generated formulas are

 $\forall A \text{ (false } \land q(A) \supset false)$ 

false ⊃false.

Both of them are trivially true.

Example 4.5.2: When a theorem includes a dominant atom with a type predicate, we have the effect to perform the usual structural inuction. Suppose we prove (cf.Kowalski [13] pp.221-222)

theorem(right-identity-of-append).

∀ A:list append(A,[],A).

Note the dominant atom list(A). Because each definite clause is inducible with 0, we can apply the induction immediately and two generated formulas are

```
append([],[],[]),
\forall U,A (append(A,[],A) \supset append([U|A],[],[U|A])).
```

Example 4.5.3: Theorems may include existentially quantified variables. Suppose we prove theorem(last-can-be-first-of-reversed).

 $\forall A,V,C \text{ (append}(A,[V],C) \supset \exists B \text{ reverse}(C,[V|B])).$ 

Note the dominant atom append(A, [V], C). Because each definite clause is inducible with 0, we can apply the induction immediately and two generated formulas are

 $\forall V \exists B \text{ reverse}([V],[V|B]),$ 

 $\forall U,A,V,C (\exists B \text{ reverse}(C,[V|B]) \supset \exists B \text{ reverse}([U|C],[V|B])).$ 

Note that the generated subgoals still remain within S-formulas.

# 5. Manipulation of Computational Induction Schemes

### 5.1. Mergible Schemes

In section 4,we cited examples with only one dominant atom intensionally. But sometimes more than two different induction schemes are suggested from theorems to be proved.

Example 5.1: Suppose we prove the second formula generated from reverse-reverse theorem(first-last).

```
\forall A,B,C,U \; (reverse(C,A) \; \land append(C,[U],B) \; \supset reverse(B,[U|A])).
```

Then both reverse(C, A) and append(C, [U], B) are dominant. Let  $Q_1$  and  $Q_2$  be  $Q_1(A,B,C,U)$ : append $(C,[U],B) \supset reverse(B,[U|A])$ ,

 $Q_2(A,B,C,U)$ : reverse(C,A)  $\supset$ reverse(B,[U|A]).

Then reverse(C, A) and append(C, [U], B) suggests two different induction schemes.

$$\forall B, U \ Q_1([:],B,[:],U) \ \forall A,B,C,U,A_1,V, \ (Q_1(A_1,B,C,U) \land append(A_1,[V],A) \supset Q_1(A,B,[V|C],U))$$

$$\forall A,B,C,U \ (reverse(C,A) \land append(C,[U],B) \supset reverse(B,[U|A]))$$

$$\forall A,U \ Q_2(A,[U],[:],U) \ \forall A,B,C,U,V \ (Q_2(A,B,C,U) \supset Q_2(A,[V|B],[V|C],U))$$

$$\forall A,B,C,U \ (reverse(C,A) \land append(C,[U],B) \supset reverse(B,[U|A]))$$

In the example above, the substitution to N in each scheme coincides and it suggests a possibility to apply both inductions simultaneously. In order to justify such a manipulation, we describe an equivalence preserving transformation in the next section.

#### 5.2. Tamaki-Sato's Transformation

The Tamaki-Sato's transformation system is developed for Prolog programs based on unfold/fold transformations. The entire process proceeds as follows [16].

#### Transformation Process

```
P_0 := the initial program; D_0 := \{\}; mark every clause in P_0 "foldable"; for i := 1 to arbitrary N apply any of the transformation rules to obtain P_i and D_i from P_{i-1} and D_{i-1};
```

Example 5.2.1: Before starting the transformation, the initial program is given, e.g.,

```
\begin{array}{c} P_0: C_1. \ append([\ ],L,L). \\ C_2. \ append([X|L],M,[X|N]):= append(L,M,N). \\ C_3. \ reverse([\ ],.[\ ]). \\ C_4. \ reverse([X|L],M):= reverse(L,N),append(N,[X],M). \\ and \ D_0 \ is \ initialized \ to \ \{\}. \end{array}
```

The basic part of Tamaki-Sato's transformation system consists of three transformation rules, i.e., definition, unfolding and folding.

Definition: Let C be a clause of the form  $p(X_1, X_2, ..., X_n) = A_1, A_2, ..., A_m$  where

- (a) p is an arbitrary predicate appearing neither in P<sub>i-1</sub> nor in D<sub>i-1</sub>
- (b) X<sub>1</sub>, X<sub>2</sub>,..., X<sub>n</sub> are distinct variables, and
- (c)  $A_1, A_2, \ldots, A_m$  are atoms whose predicates all appears in  $P_0$ . Then let  $P_i$  be  $P_{i-1} \cup \{C\}$  and  $D_i$  be  $D_{i-1} \cup \{C\}$ . Do not mark C "foldable".

Example 5.2.2: Suppose we need the conjunction of atoms each induction scheme is accounting for in example 5.1. Then we introduce it by the following definition.

```
C_5. new-p(L,M,N,X) :- reverse(N,L),append(N,[X],M).
Then P_1 = \{\underline{C_1},\underline{C_2},\underline{C_3},\underline{C_4},C_5\} and D_1 = \{\underline{C_5}\}. The underlines indicate "foldable" clauses.
```

Unfolding: Let C be a clause in  $P_{i-1}$ , A be an atom in its body and  $C_1, C_2, \ldots, C_k$  be all the clauses in  $P_{i-1}$  whose heads are unifiable with A. Let  $C'_i$  be the result of resolving C with  $C_i$  on A. Then let  $P_i$  be  $\{P_{i-1} - \{C\}\} \cup \{C'_1, C'_2, \ldots, C'_k\}$  and  $D_i$  be  $D_{i-1}$ . Mark each  $C'_i$  "foldable" unless it is already in  $P_{i-1}$ .

Example 5.2.3: When  $C_5$  is unfolded at its first atom in the body, we obtain  $P_2 = \{C_1, C_2, C_3, C_4, C_5\}$  $C_4, C_6, C_7$  and  $D_2 = \{C_5\}$  where

 $\overline{C_6}$ . new-p([],M,[],X) :- append([],[X],M).

 $C_7. \ \ \text{new-p}(L,M,[Y|N],X) := \text{reverse}(N,L_1), \\ \text{append}(L_1,[Y],L), \\ \text{append}([Y|N],[X],M).$  $C_6$  and  $C_7$  are still unfoldable into

 $C_7'. \ \ \text{new-p}(L,[Y|M],[Y|N],X) := \text{reverse}(N,L_1), \\ \text{append}(L_1,[Y],L), \\ \text{append}(N,[X],M).$ C'6. new-p([],[X],[],X). and we get  $P_3 = \{\underline{C_1}, \underline{C_2}, \underline{C_3}, \underline{C_4}, \underline{C_6}, \underline{C_7}\}$  and  $D_3 = \{C_5\}$ .

Folding: Let C be a clause in  $P_{i-1}$  of the form  $A_0:=A_1,A_2,\ldots,A_n$  and  $C_{folder}$  be a clause in  $D_{i-1}$  of the form  $B_0: B_1, B_2, \ldots, B_m$ . Suppose there is a substitution  $\sigma$  and a subset  $\{A_{i_1},A_{i_2},\ldots,A_{i_m}\}$  of the body of C such that the following conditions hold.

- (b)  $\sigma$  substitutes distinct variables for the internal variables of  $C_{folder}$  and moreover those variables occur neither in A nor in  $\{A_1,A_2,...,A_n\} = \{A_{i_1},A_{i_2},...,A_{i_m}\}$ , and

(c) C is marked "foldable" or m < n. Then let  $P_i$  be  $(P_{i-1} - \{C\}) \bigcup \{C'\}$  and  $D_i$  be  $D_{i-1}$  where C' is a clause with head A and body  $(\{A_1,A_2,\ldots,A_n\}-\{A_{i_1},A_{i_2},\ldots,A_{i_m}\})\bigcup\{\sigma(B)\}$ . Let C inherit the mark of C.

Example 5.2.4: By folding the whole body of  $C_7$  by  $C_5$ , we obtain  $P_4 = \{\underline{C_1}, \underline{C_2}, \underline{C_3}, \underline{C_4}, \underline{C_6}, \underline{C_8}\}$ and  $D_4 = \{C_5\}$  where

 $C_8. \ \text{new-p}(L,[Y|M],[Y|N],X) := \text{new-p}(L_1,M,N,X), \\ \text{append}(L_1,[Y],L).$ 

The most important property of this transformation system is stated as follows [16].

 $P_N$  is equivalent to  $P_0 \bigcup D_N$  in the minimum Herbrand model semantics,i.e., the min-Equivalence Preservation Theorem imum Herbrand models of them are identical.

Example 5.2.5: Through the previous transformation process, we reach a program

 $P_4$ :  $C_1$ . append([],L,L).

 $C_2$ . append([X|L],M,[X|N]) := append(L,M,N).

C3. reverse([],[]).

 $C_{4}.\ reverse([X|L],M): \ reverse(L,N), append(N,[X],M).$ 

 $C_6'$ . new-p([],[X],[],X).

 $C_8. \ \text{new-p}(L,[Y|M],[Y|N],X) := \text{new-p}(L_1,M,N,X), \\ \text{append}(L_1,[Y],L).$ 

which is equivalent to

 $P_0 : C_1$ . append([],L,L).

 $C_2$ . append([X|L],M,[X|N]) :- append(L,M,N).

C3. reverse([],[]).

 $C_4$ . reverse([X|L],M) : reverse(L,N),append(N,[X],M).

 $D_4:C_5$ . new-p(L,M,N,X) := reverse(N,L),append(N,[X],M).

# 5.3. Derivation of Merged Schemes

We keep each induction scheme in a triple  $(A,Q,\{\sigma_i(C_i)\})$ , i.e.,an atom A accounting for, a goal formula Q and an instanciated program  $\{\sigma_i(C_i)\}$ .

Example 5.3.1: Suppose we are trying to prove the theorem first-last. Before merging, we have two induction schemes I and J whose instanciated programs are

I : reverse([],[]).

```
reverse([X|L],M) := reverse(L,N), append(N,[X],M).
J : append([],[X],[X]).
append([X|L],[Y],[X|M]) := append(L,[Y],M).
```

An induction scheme J accounting for  $q(s_1, s_2, ..., s_m)$  is said to be mergible to an induction scheme I accounting for  $p(t_1, t_2, ..., t_n)$  (cf. [3] pp.191-194) when

- (a)  $p(t_1, t_2, ..., t_n)$  and  $q(s_1, s_2, ..., s_m)$  share at least one free variable,
- (b) Every recursive definite clause of I is mergible to only one definite clause of J and
- (c) No two recursive definite clauses of I is mergible to a same definite clause of J.

where a recursive definite clause  $A_0 := A_1, A_2, \ldots, A_{\alpha}$  is said to be mergible to a recursive definite clause  $B_0 := B_1, B_2, \ldots, B_{\beta}$  when the following conditions are satisfied. Let  $A_0$  be  $\tau_0(p(t_1, t_2, \ldots, t_n))$  and  $B_0$  be  $= \tau'_0(q(s_1, s_2, \ldots, s_m))$ .

- (a)  $\tau_0$  and  $\tau_0'$  substitute an identical term to each common variable of  $p(t_1, t_2, ..., t_n)$  and  $q(s_1, s_2, ..., s_m)$ ,
- (b) There is a one-to-one correspondence between recursive calls in the bodies such that the corresponding  $r_i$  and  $r'_j$  substitute an identical term to each common variable of  $p(t_1, t_2, ..., t_n)$  and  $q(s_1, s_2, ..., s_m)$ .

Example 5.3.2: The previous two induction schemes are ovbiously mergible, because for the second clauses,

- (a) When reverse(C, A) is unified with reverse([X|L], M) by  $r_0 = < C \leftarrow [X|L], A \leftarrow M >$ , append(C, [U], B) is unifiable with append([X|L], [Y], [X|N]) by  $r'_0 = < C \leftarrow [X|L], B \leftarrow [X|N], U \leftarrow Y >$
- (b) There is only one recursive calls reverse(L, N) and append(L, [Y], N) in each second definite clause and they are instances of reverse(C, A) by  $\tau_1 = \langle C \Leftarrow L, A \Leftarrow N \rangle$  and of append(C, [U], B) by  $\tau_1 = \langle C \Leftarrow L, B \Leftarrow N, U \Leftarrow Y \rangle$ .

The mergibility guarantees that a transformation sequence like one stated in example 5.2.1—5 is applicable. Note that the transformation is a routine and we can apply it mechanically. When two schemes are mergible, we derive a new scheme as follows.

- (a) Define  $new-p(X_1, X_2, ..., X_l)$  by the conjunction of  $p(t_1, t_2, ..., t_n)$  and  $q(s_1, s_2, ..., s_m)$ .
- (b) Unfold at  $p(t_1, t_2, ..., t_n)$  and  $q(s_1, s_2, ..., s_m)$  once in the bodies of definite clauses.
- (c) Fold if possible.

Then the obtained definite clauses represent a new scheme with  $new-p(X_1, X_2, ..., X_l)$  and  $Q = G_{p(i_1, i_2, ..., i_n), q(s_1, s_2, ..., q_m)}[true, true] \downarrow$ .

Example 5.3.3: As was shown in example 5.2.1—5, we can obtain a definite clause program for new-p from the previous two induction schemes

```
\begin{array}{l} \text{new-p([\ ],[X],[\ ],X).} \\ \text{new-p(L,[Y]M],[Y]N],X) := \text{new-p(L_1,M,N,X),append(L_1,[Y],L).} \end{array}
```

and the theorem to be proved is now

 $\forall$  A,B,C,U (new-p(A,B,C,U) $\supset$ reverse(B,[U|A])).

Hence the new scheme accounting for new-p represents

```
\forall U \ reverse([U],[U]) \quad \forall A,B,C,U,A_1,V \ (reverse(B,[U|A_1]) \ \land append(A_1,[V],A) \ \supset reverse([V|B],[U|A]))
\forall A,B,C,U \ (reverse(C,A) \ \land append(C,[U],B) \ \supset reverse(B,[U|A]))
```

Merging of computational induction schemes sometimes results in surprisingly simple schemes.

```
Example 5.3.4: Suppose we prove a theorem
     theorem(equivalence-of-flatten-and-mc-flatten)
         \forall W,A,B,C \text{ (flatten(W,A) } \land mc\text{-flatten(W,B,C)} \supset append(A,B,C)).
where flatten and mc-flatten are procedures collecting leaves of S-expressions defined as
follows [3]. (mc-flatten(W, [], A) performs the same task as flatten(W, A) more efficiently.)
      flatten(X,[X]) := atom(X)
      flatten([X,Y],L) := flatten(X,L_1), flatten(Y,L_2), append(L_1,L_2,L).
      mc-flatten(X,M,[X|M]):- atom(X).
      mc-flatten([X|Y],M,N) :- mc-flatten(Y,M,L),mc-flatten(X,L,N).
Then by defining a new predicate by
      \mathtt{new-p}(Z,L,M,N) \ : \ \mathtt{flatten}(Z,L),\mathtt{mc-flatten}(Z,M,N).
and applying the transformations, we obtain a new definition
      new-p(X,[X],M,[X|M]) := atom(X).
      \mathtt{new-p}([X|Y],L,M,N) := \mathtt{append}(L_1,L_2,L), \mathtt{new-p}(X,L_1,L_3,N), \mathtt{new-p}(Y,L_2,M,L_3).
 The scheme accounting for new-p(W, A, B, C) is
                                 \forall U,B (atom(U) \supset append([U],B,[U|B]))
\forall A,B,C,A_1,A_2,A_3 \; (\text{append}(A_1,A_2,A) \; \land \text{append}(A_1,A_3,C) \; \land \text{append}(A_2,B,A_3) \; \supset \text{append}(A,B,C))
                  \forall W,A,B,C \text{ (flatten}(W,A) \land mc\text{-flatten}(W,B,C) \supset append}(A,B,C))
```

The second formula is the associativity of append for  $A_1,A_2$  and B.

#### 6. Discussions

Computational induction is due to de Bakker and Scott [2]. Its use in mechanical verification is investigated by Weyrauch and Milner [18] and Gordon etc [8]. The simplest form of its use in Prolog was pointed out by Clark [5]. (Integration of induction into a logic programming system as an inference rule is investigated by Hagiya and Sakurai [9].) But our use is more general in the sense that atoms accounted for need not be in general form  $p(X_1, X_2, ..., X_n)$ , e.g. Example 4.4.5 and 4.5.3. Our method to merge computational induction schemes is new as far as we know.

In our verification system, if we can apply computational inductions, we prefer them. But when we can't because of the conditions and heuristic constraint, we resort to well-founded induction. Formulation of well-founded induction is done similarly with BMTP-like heuristics [12].

#### 7. Conclusions

We have shown how to formulate computational induction formulas in verification of Prolog programs and its advantages. Compared to BMTF, use of computational induction (a) need not guarantee termination by any well-founded ordering, (b) does not require the restriction that procedures be terminating, (c) generate induction goals 1 less than BMTP-like well-founded inductions, (d) generate simpler induction formulas more first order inferences are already performed to and (c) accompodate naive structural inductions and well-founded inductions in many cases. This method to formulate induction formulas is an element of our verification system for Prolog programs under development.

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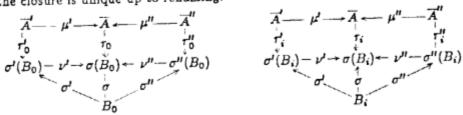
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## Appendix. Closure of Atom

Theorem Closure is unique up to renaming.

Proof: Suppose A has two closures  $\overrightarrow{A}$  and  $\overrightarrow{A}''$ . Then from the condition (b), they are unifiable. Let its most general instance be  $\overrightarrow{A} = \mu'(\overrightarrow{A}) = \mu''(\overrightarrow{A}'')$ . Suppose a head of a recursive definite clause  $B_0$  is unifiable with  $\overrightarrow{A}$  by an m.g.u.  $\tau_0 \circ \sigma$ . Hence  $(\tau_0 \circ \mu') \circ \sigma$  is a unifier of  $\overrightarrow{A}'$  and  $B_0$  and  $(\tau \circ \mu'') \circ \sigma$  is a unifier of  $\overrightarrow{A}''$  and  $B_0$ . Because  $\overrightarrow{A}$  and  $\overrightarrow{A}''$  are closures of A,  $B_0$  is unifiable with  $\overrightarrow{A}'$  by an m.g.u.  $\tau'_0 \circ \sigma'$  and unifiable with  $\overrightarrow{A}''$  by an m.g.u.  $\tau''_0 \circ \sigma''$ . This means that for some  $\nu'$  and  $\nu''$ ,  $\sigma = \nu' \circ \sigma' = \nu'' \circ \sigma''$ . For all i such that  $B_i$  is a recursive call,  $\sigma(B_i) = \nu' \circ \sigma'(\overrightarrow{A}) = \nu' \circ \tau'_i(\overrightarrow{A}) = \nu'' \circ \sigma''(\overrightarrow{A}'') = \nu'' \circ \tau'_i(\overrightarrow{A}'')$ . Hence  $\sigma(B_i)$  is a common instance of  $\overrightarrow{A}'$  and  $\overrightarrow{A}''$ . Because  $\mu' \circ \mu''$  is an m.g.u. of  $\overrightarrow{A}'$  and  $\overrightarrow{A}''$ , there exists a substitution  $\tau_i$  such that  $\sigma(B_i) = \tau_i \circ \mu'(\overrightarrow{A}') = \tau_i \circ \mu''(\overrightarrow{A}'')$ . Then  $\tau_i$  satisfies the condition of the closedness and  $\overrightarrow{A}$  is a closure of A. Because of the condition (c),  $\overrightarrow{A}'$  and  $\overrightarrow{A}''$  are variants. Hence the closure is unique up to renaming.



Computation of  $p(t_1, t_2, ..., t_n)$ 

```
i:=-1; A_0:=p(t_1,t_2,\ldots,t_n); P_0:= the set of recursive definite clauses defining p;
repeat
   i := i + 1; select a recursive definite clause C in P_i fairly;
    if the head of C is unifiable with A_i
   then A_{i+1} := closurel(A_i, C); P_{i+1} := P_i - \{C\};
    else A_{i+1} := A_i; P_{i+1} := P_i;
until all heads of definite clauses in P_i are not unifiable with A_i
return A_{i+1}
closure1(A, "B<sub>0</sub> := B<sub>1</sub>, B<sub>2</sub>, . . . , B<sub>m</sub>");
i := -1; A_0 := A;
repeat forever
    i := i + 1;
    let \tau_0 \circ \sigma be an m.g.u. of A_i and B_0
    where \tau_0 and \sigma are the restrictions to A_i and B_0;
    let B_0', B_1', B_2', \ldots, B_1' be variants of atoms with p in \sigma(B_0 : B_1, B_2, \ldots, B_m)
    without shared variable by an appropriate renaming;
    B := \text{most specific common generalization of } B_0^t, B_1^t, B_2^t, \dots, B_1^t;
    if B is an instance of A_i then return A_i else A_{i+1} := B_i
```

where most specific common generalization is the dual of most general common instanciation, i.e., E is a most specific common generalization of  $E_1, E_2, ..., E_l$  when

- (a) E<sub>1</sub>, E<sub>2</sub>,..., E<sub>l</sub> are instances of E.
- (b) E is an instance of any E' satisfying (a).

It is easily obtained by comparing the corresponding subexpressions.