

# A simple algorithm to construct a consistent extension of a partially oriented graph

Dorit Dor

Computer Science Department  
School of Mathematical Sciences  
Tel-Aviv University  
69978  
Israel

Michael Tarsi \*

Cognitive Systems Laboratory  
Computer Science Department  
University of California, Los Angeles, CA 90024

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## Abstract

A *Partially directed acyclic graph*, (*pdag*), is a graph which contains both directed and undirected edges, with no directed cycle in its directed subgraph. An oriented extension of a *pdag*  $G$  is a fully directed acyclic graph (*dag*) on the same underlying set of edges, with the same orientation on the directed subgraph of  $G$  and the same set of *vee-structures*. A *vee-structure* is formed by two edges, directed toward a common head, while their tails are nonadjacent. A simple polynomial-time algorithm is presented, to solve the following problem: *Given a pdag, does it admit an oriented extension?* The problem was stated by Verma and Pearl, while studying the existence of causal explanation to a given set of observed independencies.

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\*Visiting professor. Permanent address: Computer Science Department, School of Mathematical Sciences, Tel-Aviv University, 69978, Israel. Supported in part by the Air Force Office, AFOSR 90 0136 and by Northrop MICRO, PO 11510.

# 1 Introduction

A *Partially directed acyclic graph*, (*pdag*), is a graph which contains both directed and undirected edges, with no directed cycle in its directed subgraph. An oriented extension of a pdag  $G$  is a fully directed acyclic graph (dag) on the same underlying set of edges, with the same orientation on the directed subgraph of  $G$  and the same set of *vee-structures*. A vee-structure is formed by two edges, directed toward a common head, while their tails are nonadjacent. These definitions as well as some background and motivation are stated and explained in [1]. While studying the existence of causal explanation to a given set of observed independencies, Verma and Pearl [1] have faced the following combinatorial problem, to which we refer here as "PDX" (PDag eXtensibility): *Given a pdag, does it admit an oriented extension?*

In Section 2 of [1] the authors present an algorithm for PDX, which is conjectured, however not proven, to be polynomial. Another algorithm, given by Verma in [3], although it runs in linear time, is rather complicated and less intuitive. We present here a simple polynomial-time algorithm to solve the above problem.

## 2 The algorithm

Our algorithm selects first a vertex  $x$  to be the sink of the extension and recursively proceed to the subgraph obtained by the removal of the sink and all edges incident to it:

Algorithm extend( $G$ : pdag);

begin (extend)

$G' := G$ ;  $A := G$ ;

while  $A$  is not empty do

begin (iteration)

Select a vertex  $x$  which satisfies the following properties in the subgraph  $A$ :

- a.  $x$  is a sink (no edge  $(x, y)$  in  $A$  is directed outward from  $x$ )
- b. For every vertex  $y$ , adjacent to  $x$ , with  $(x, y)$  undirected,  $y$  is adjacent to all the other vertices which are adjacent to  $x$ ;

If such  $x$  is not found, then the algorithm stops and returns a negative answer ( $G$  does not admit any extension);

If  $x$  is found, let all the edges which are incident to  $x$  in  $A$  be directed toward  $x$  in  $G'$  ( $G'$  is meant to form the output);

$A := A - x$  (remove  $x$  and all the edges incident to  $x$ )

end (iteration);

return  $G'$  (an extension of the input pdag  $G$ )

end (extend).

### 3 Validity and complexity

An extension of  $G$ , if it exists, is a dag and as such it contains a sink. To become a sink of the extension  $G'$  a vertex  $x$  must satisfy property  $a$ . (of the iteration phase above) in  $G$ . To avoid the creation of new vee-structures, while directing all edges toward  $x$ , it should also satisfy property  $b$ . Hence a vertex which satisfies both properties  $a$ . and  $b$ . is indeed necessary for the existence of an extension. To justify our recursive method we should show first that the removal of a sink  $x$  from the extension  $G'$  provides an extension  $G' - x$  of the pdag  $G - x$ , obtained when the same vertex is removed from the input pdag  $G$ : No directed cycle can be formed by the removal of  $x$  and hence  $G' - x$  is still a dag. In the general case a new v-structure might be generated by the removal of edges if a single edge is deleted from a triangle, whose other two edges now form a v-structure. In the case on hand, however, all the edges incident to  $x$  are deleted, thus, the number of edges removed from any triangle is either 0 or 2.  $G' - x$  is hence indeed an extension of  $G - x$ . To complete the proof we should notice that no existing extension is missed by selecting the specific vertex  $x$  to be a sink. If there exists an extension where  $x$  is not a sink then it will still remain an extension if the edges going out of  $x$  are reoriented toward  $x$ . All redirected edges are incident to  $x$  and since  $x$  is now a sink no directed cycle was formed. Also no new v-structures are created due to property  $b$ . of the selected vertex  $x$ .

For the complexity analysis note that there are  $|V|$  iterations where every edge is searched at most twice (once for each endvertex). The time complexity is thus  $O(|V||E|)$ .

### 4 Some concluding remarks

A dag with no vee-structure is chordal (any orientation of a chordless cycle contains a vee-structure). Consequently, if  $G$  contains no vee-structure then its underlying graph should be chordal. A known characterization of chordal graphs states that all such graphs can be constructed, starting with an isolated vertex, by successive insertion of new vertices, each adjacent to a clique in the existing graph. When our algorithm is applied to a graph  $G$  with no oriented edges then property  $b$ . states that the neighbors of  $x$  form a clique. In this case  $G$  admits an extension if and only if it is chordal. Our algorithm is a natural generalization of a naive chordality test, based on the above characterization, to the case where v-structures are allowed, but no new ones should be formed. Chordality can be tested in linear time [2] and hence it takes linear time to test the existence of an extension where the input has no vee-structures. We believe that linear-time chordality algorithm can be modified to a general linear-time algorithm for *PDX*.

## References

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