New Shape Analysis Techniques for Automatic Parallelization of C Codes

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New shape analysis techniques for automatic parallelization of C codes

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Abstract

Automatic parallelization of codes with complex data structures is becoming very important. These complex, and often, recursive data structures are widely used in scientific computing. Shape analysis is one of the key steps in the automatic parallelization of such codes. In this paper we extend the Static Shape Graph (SSG) method to enable the successful and accurate detection of complex doubly linked structures. In addition, these techniques have been implemented in a compiler, which has been validated for several C codes. In particular, we present the results the compiler achieves for the C sparse LU factorization algorithm. The output SSG for this case study perfectly describes the complex data structure used during the LU code.

1 Introduction

Regarding high performance computing, it is clear that compilers represent a key tool that should take care of optimizing the applications that must be executed efficiently in parallel computers. A good deal of work has been done in the area of array dependence analysis [2] with notable success. However, non-numerical and numerical applications based on complex and dynamic data structures are becoming more and more widely used lately. These complex, and often, recursive data structures are based on dynamic allocation and are generated at execution time. Parallelization of an application requires the compiler's special knowledge about the underlying semantic of the data structure. With these assumptions, shape analysis becomes a first step in the data dependence test for such kinds of codes. The aim of this phase is to find out at compile time the shape of the heap.

In this work we present some important modifications to the shape analysis method developed by Sagiv et al. [12]. In addition, we check our improvements with a real numerical application like the non-symmetric sparse LU decomposition based on a one-dimensional doubly linked list.

The organization of the paper follows. In the next section we revise the approaches currently available to solve shape analysis problems. Section three presents the motivating example. Our shape analysis techniques are presented in section four. Conclusions and future work close the paper.

2 Related Work

There are several methods addressing the shape analysis problem. Some of these are based on explicit annotations, as in Hummel et al. [7]. These methods are based on programmer annotations describing the data structure.

Other approximations are based on access paths. Henderson et al. [5] use “path matrix analysis” that contains “access paths” between pointers. Matsumoto et al. [10] use “normalized” path expressions to maintain the “alias-pair” between pointers. These methods cannot handle cyclic structures like double linked lists and trees with parent pointers. Finally, there are methods based on graphs. In the graph, the nodes represent “storage chunks”, and the edges references between them.

One of the first relevant works on this topic was developed by Jones et al. [8]. In this work, the authors focus on the shape analysis of programs with destructive updating. They bind, to each program point, a set of graphs which describe all potential alias relationships that can arise at execution time. In addition, they use a “k-limited” approximation in which all nodes beyond a k selectors path are joined in a summary node. Horwitz et al. [6] presented another variation on k-limited graphs, called “storage graphs”. Also, the authors maintain a set of storage graphs at each program point. The main drawbacks of these methods are: (1) the number of shape graphs that can arise for each program point is very high, leading to a great deal of computational and memory overhead; and (2) the node analysis beyond the “k-limit” is very inexact.

On the other hand, there are approximations in which each program point has an associated graph which covers all possible shape graphs combinations, instead of all these possible graphs independently [9, 3, 11, 12]. The result of joining all the information, previously represented by different shape graphs, in a single one, is a lack of accuracy in the representation, but on the other hand, it leads to a practical shape analysis algorithm. Larus et al. [9], use a variation of “k-limited” graphs called “alias graphs”, and introduce
summary nodes using "s-l limiting". This method works well only for simple data structures like trees and lists. It is expensive by its complex meet, node summary and node labeling operations.

The algorithm presented by Chase et al. [3] is not "k-limited". Their abstraction "Storage Shape Graph" contains one node for each variable and one for each location site in the program. In this method, the count of references from the heap to this node (0, 1, inf), is stored for each node in the graph. This way, the k-limited drawback is avoided. This algorithm is able to detect a single linked list even when new elements are appended to the end of the list. However, it is not powerful enough to detect insertions of elements in the middle of the list.

Plevyak et al. [11] work is based on the Chase’s method. They extend the previous "Storage Shape Graph" into the "Abstract Storage Graph" (ASG) in order to solve the main problems arising in the first one. However, in the same way as Chase’s method, their comparison and compression operations are complex and expensive.

The method presented by Sagiv et al. [12] is based on what they call "Static Shape Graphs" (SSG). The main difference between this method and previous ones lies in the node-name scheme they use for the nodes. Their graph contains nodes only for heap locations pointed to by program variables. Some very interesting properties are: (1) alias relationships are easier to find; (2) the determinism is better preserved in the graph due to they always carry out a "strong nullification" equivalent to "strong update" (substitution of a reference by another one, without keeping the previous one).

SSG the union and comparison operations are very simple due to this node-naming scheme.

However this method cannot analyze doubly linked structures which are widely used in C codes, like Sparse LU factorization. In this case the SSG is not able to accurately represent the data structure of the Sparse LU factorization. Each column of the sparse matrix is represented by a doubly linked list. In addition, a different doubly linked list is needed to point to the first element of each list (column).

We propose combining the Sagiv’s method [12] and the Abstract Storage Graph (ASG) proposed by Plevyak et al. [11] to achieve a more precise shape analysis of this type of structures which will allow us to automatically parallelization of C codes with complex data structures. The extended SSG proposed in this work introduces two main modifications:

- There will be several summary nodes in the SSG, allowing us to summarize different structure and type elements into different summary nodes, each one with its own properties.
- The “shared" attribute is assigned to each selector and to pairs of selectors in cycle links. With this modification we achieve a more accurate representation of the doubly linked structures.

3 Motivating example: sparse LU factorization

The kernel of many computer-assisted scientific applications is to solve large sparse linear systems. We find examples of these kind of applications in optimization problems, linear programming, simulation, circuit analysis, fluid dynamic computation, and numeric solutions of differential equations in general.

Furthermore, this problem presents a good case study and is a representative computational code for many other irregular problems. Actually, this problem represents those in which the computational load grows with the execution time (fill-in) and matrix coefficients change their coordinates due to row/column permutations (pivoting).

More precisely, our working example application solves non-symmetric sparse linear systems by applying the LU factorization of the sparse matrix, computed by using a general method [1, 4]. These methods directly solve the sparse problem and share the same loop structure of the corresponding dense code (the one we see in Fig. 1).

In this Fig. 1, we show an in-place code for the direct right-looking LU algorithm, where an n-by-n matrix A is factorized. The code includes a row pivoting operation (partial pivoting) to provide numerical stability and preserve sparsity.

Usually, in order to save both memory and computation overhead, zero entries of sparse matrices are not explicitly stored. A wide range of methods for storing the nonzero entries of sparse matrices have been developed [4]. Here, we will consider only linked list data structures.

The partial-pivoting LU decomposition stores the coefficient matrix in a one-dimensional doubly linked list (see Fig. 2 (b)), to facilitate the insertion of new entries and to allow column permutations.

Analyzing the sparse LU algorithm with Sagiv’s method, the resulting SSG is shown in Fig. 3. Here, we can see that the same summary node, "s_0", refers to both the elements belonging to the header list and the ones in the column linked lists. This way it is impossible to discern between the two different data structures (header and columns).

Furthermore, the summary node is shared, (is = true), which means that the summarized nodes are referenced from the heap more than once, when actually, they are referenced...
4 A modified shape analysis algorithm

This section focuses on the description of the new techniques which solve the previous problem. However, before this it is necessary to briefly introduce the SSG notation used in [12].

4.1 SSG Notation.

An SSG is a finite, labeled, directed graph that approximates the actual stores that can arise during program execution. The shape-analysis algorithm itself is an iterative procedure that computes an SSG at every program point.

An SSG $SG$ consists of two kinds of nodes, variables ($PVar$) and shape-nodes, and two kinds of edges, variable-edges and selector-edges. An SSG is represented by a pair of edges sets, $<E^v,E^s>$, where:

- $E^v$ is the graph's set of variable-edges, each of which is denoted by a pair of the form $[x,n]$, where $x \in PVar$ and $n$ is a shape-node.
- $E^s$ is the graph's set of selector-edges, each of which is denoted by a triple of the form $<s,sel,t>$ where $s$ and $t$ are shape-nodes, and $sel$ is a selector.

Shape-nodes are named using a (possibly empty) set of pointer variables, $X$. The set $shape\_node \; s(SG^s)$ is a subset of $\{n_x \mid X \subseteq PVar\}$. A shape-node $n_X$, where $X \neq \emptyset$, represents the cons-cell (storage chunk) pointed to by exactly the pointer variables in the set $X$, in any given concrete store.

The shape-node $n_\emptyset$ (summary node) can represent multiple cons-cells of a single concrete store.

Each shape-node $n$ in an SSG has an associated Boolean flag, denoted by $is\_n(n)$ (is shared). When $is\_n(n)=true$, indicates that the cons-cells represented by $n$ may be the target of pointers emanating from two or more distinct cons-cells fields. On the other hand, $is\_n(n)=false$ means that, if several selector edges in an SSG point to $n$, they represent concrete edges that never point to the same cons-cell in any concrete store. The function $is\_n$ is therefore of type $shape\_nodes(SG^s) \rightarrow \{false,true\}$.

Two different shape-nodes $n_x$ and $n_y$, such that $X \neq Y$ and $X \cap Y \neq \emptyset$, represent incompatible configurations of variables. Thus, for all $<n_x,sel,n_y> \in E^s$, either $X = Y$ or $X \cap Y = \emptyset$. The function compatible$^s(n_x,\ldots,n_y)$ means $\forall i,j : Z_i = Z_j \vee Z_i \cap Z_j = \emptyset$.

The "Abstract Interpretation" presents the modifications on a SSG for the six kind of statements that manipulate pointer variables $(x := nil, x.sel := nil, x := new, x := y, x.sel := y$ and $x := y.sel)$. $E^v$, $E^s$ and $is\#s$ are $E^v$, $E^s$ and $is\#$ after statement execution.

With all these definitions, we can move on to the main part of this section: the description of the new techniques we propose.

4.2 Several Summary Nodes

The SSG method [12] can contain only one summary node: the one which represents the whole storage chunk in a certain program point which is not referenced directly by any variable. However, to improve the data structure representation in many cases, we allow the existence of more than one summary node. More precisely, in our SSG there may be a summary node for each pointer type and connected component, as we describe now.

4.2.1 A Summary Node per pointer type.

If the method is constrained to a single summary node, then nodes of different structure type may be summarized in a single node. This way, all of these nodes will have the same "is shared", $is\#$, attribute. Obviously, $is\#s$ may turn to be true at a certain program point, but this is less likely to happen now that the summary nodes are actually representing less nodes.

For instance, by allowing only one summary node, two different structures, like the ones we see in figure Fig. 4 (a), are going to be represented by the same summary node, Fig. 4 (b). This way, even when only one of the structures has several references to the same node, $is\#$ becomes true for the whole summary node. Therefore, there is no way to know which structure (or if both of them) is actually sharing elements. This can be solved allowing a summary node for each different type of structure, as we can see in Fig. 4 (c). More precisely, we will consider that two structures have different type if the pointers pointing to elements of them have different type in the pointer declaration.

In order to do this, apart from the $is\#$ attribute, we associate to each node the type information ($type\#s$). For each pointer variable we keep its type ($type\_var$), which is taken from the declaratory part. With all these assumptions, the abstract semantic of the following statements should be modified:

1. Statement $[x := new]$  
   The $type\#$ of the new node $(n_{x_1})$ is set to the $type\#$ of the variable that points to it.  
   $$ type\#(n_{x_1}) = type\_var(x) $$

2. Statement $[x := y]$  
   All the nodes preserve their type, and the new nodes (now referenced by "x") take the type of the nodes pointed to by "y".  
   $$ type\#(n_x) = type\#(n_{x_{-x_1}}) $$

3. Statement $[x := y.sel]$  
   A node materialization takes place. The type of the new node is the same as the type of the node from which it is materialized.
4. The summarization of nodes not directly referenced by pointer variables, varies as well. Now, only nodes of the same type and not pointed to by any variable can be summarized.

5. During node matching for the union or comparison of graphs, apart from the set of variables referenced by the node, it is now also necessary to match their type. This only affects the summary nodes since the others will have the type equal to that of the variables which reference them.

### 4.2.2 A Summary Node per connected component.

With the previous modifications we can maintain, for each graph, different summary structures for different "types" of structures. Now, if we deal with several structures of the same type, the corresponding nodes not directly pointed to by variables, will be summarized in a single summary node. Again, it may be of importance to explicitly distinguish these structures, even when they have the same "type". For instance, in Fig. 5 (a) we can see to different structures which do not share any element. However, the method proposed in [12] leads to the SSG presented in Fig. 5 (b). Since there is a single summary node, there is no way to detect that one of the structures should have the "is shared" attribute set to false.

To solve this problem and let the method reach a SSG like the one presented in Fig. 5 (c), each node is annotated with an additional attribute: the structure to which this node belongs, $structure#$. This $structure#$ attribute has the same value for all nodes connected by a path. More precisely, we define the set of nodes connected to a given node "n", as the set of nodes that can be found in any path toward or from node "n":

$$C[E^{#}](n) = \{ n_{j} | \exists n_{1}, \ldots, n_{i} (< n, sel_{i}, n_{1} >, \ldots, < n_{i}, sel_{i+1}, n_{j} >) \in E^{#} \}$$

Again, the abstract semantic of the following statements is modified:

1. **Statement $x := y$**

   The graph connectivity does not change for this statement. The new nodes (now pointed to by ‘x’) will have the same $structure#$ as the nodes pointed to by ‘y’.

2. **Statement $[x := nil]$**

   This statement can break a connected component creating two new ones.

   $\forall n_{x}, x \in X, < n_{x}, sel, n_{z} \in E^{#} :$

   - if $C[E^{#}](n_{x}) \cap C[E^{#}](n_{z}) = \emptyset$ then
     $\forall n \in C[E^{#}](n_{x})$, $structure^{#}(n) = new\_structure,$
     $\forall m \in C[E^{#}](n_{z})$, $structure^{#}(m) = new\_structure$
   - if $C[E^{#}](n_{x}) \cap C[E^{#}](n_{z}) \neq \emptyset$, $structure^{#}$ does not change.

When the connected components of the nodes $n_{x}$ and $n_{z}$ do not have any node in common, it is clear that we are actually dealing with two different connected components. Therefore, the $structure#$ attribute is changed for all nodes in each of them.

3. **Statement $[x,.sel := y]$**

   This statement can merge two previously unconnected components. Since any assignment to “x” or “x,.sel” is always preceded by “x := nil” or “x,.sel := nil” respectively, this statement cannot break any connection.

   $\forall n_{x}, n_{y}, [x, n_{x}],[y, n_{y}] \in E^{#}$,
   $< n_{x}, sel, n_{y} > \in E^{#}$, $compatible^{#}(n_{x}, n_{y})$:

   - $\forall n \in C[E^{#}](n_{x}), \forall m \in C[E^{#}](n_{y})$
     $structure^{#}(n) = structure^{#}(m) = new\_structure$

   The $structure#$ attribute will be equal for all nodes connected to $n_{x}$ and $n_{y}$, since now they belong to the same connected component.

4. **Statement $[x := y.sel]$**

   This statement does not break any connection in the graph. The $structure#$ attribute of the new materialized node will be the same as the one of the node from which it is materialized.

   $structure^{#}(n_{z}) = structure^{#}(n_{z_{new}})$

5. Now, during the node summarization, only those nodes not pointed to by any variable and with the same $structure#$ attribute, can be summarized in a single node.

Figure 4: (a) Real structures. (b) Shape Graph without “type”. (c) Shape Graph with “type”. In (b) List and Graph may share nodes and List may be shared.
6. Regarding the node matching, the method also takes into account that the \texttt{structure/} attributes must match as well.

4.3 Share Information per selector

In the original method, each node keeps its own \texttt{is/} attribute, which tells the compiler whether or not the node is referenced more than once from the heap. However, since this method does not take into account the selector used to reach the node, there is a potential lack of accuracy during the shape analysis. For example, Fig. 6 (a) shows a doubly linked list. Even when this list is traversed in a loop only by selector “\texttt{next}” or “\texttt{prev}”, the original method results in a single summary node with \texttt{is/} = true, Fig. 6 (b). In these kinds of situations, it is very important to keep the shared attribute for each selector, as we see in Fig. 6 (c).

Our shape analysis algorithm follows this last approach, assigning a shared attribute to each selector. Therefore, in addition to \texttt{is/} \texttt{(n)} we also introduce:

\[
\texttt{is\_sel/}(n, \texttt{sel}) \rightarrow \{\texttt{false}, \texttt{true}\}
\]

which indicates whether the node “\texttt{n}” is referenced from the heap more than once by using the selector “\texttt{sel}”. This leads to a less conservative and more accurate shape analysis for many data structures.

In order to accomplish these requirements, the abstract semantic of the following statements needs to be modified:

1. Statement \(x := \texttt{nil}\)
   The summarized nodes (no longer referenced by “\texttt{x}”) keep their \texttt{is\_sel/} attributes.
   \[
   \texttt{is\_sel/}(n_{x}, \texttt{sel}) = \texttt{is\_sel/}(n_{x}, \texttt{sel}) \lor \texttt{is\_sel/}(n_{\texttt{null}}(x), \texttt{seli}) \forall \texttt{seli}
   \]

2. Statement \(x := \texttt{new} \)
   When a new node is created, the corresponding \texttt{is\_sel/} information is initially set to “false” for all the types of selectors.
   \[
   \texttt{is\_sel/}(n_{x}, \texttt{seli}) = \texttt{false} \forall \texttt{seli}
   \]

3. Statement \(x := \texttt{y} \)
   This statement does not change \texttt{is\_sel/} attribute, since the connections in the graph are not changed.
   \[
   \texttt{is\_sel/}(n_{x}, \texttt{seli}) = \texttt{is\_sel/}(n_{\texttt{null}}(x), \texttt{seli}) \forall \texttt{seli}
   \]

4. Statement \([x, \texttt{sel} = \texttt{nil}] \)
   This statement may break references from node “\texttt{x}” by selector “\texttt{sel}”, and therefore nodes with \texttt{is\_sel/}(n, \texttt{sel}) “true” may turn to be “false”.
   To properly update the \texttt{is\_sel/} attribute, we define the following function: \texttt{is\_sel/} [\texttt{Es/}](n, \texttt{sel}) \[
   \exists n_{z1}, n_{z2}, \texttt{compatible/}(n_{z1}, n_{z2}, n) \land n_{z1}, sel, n > , n_{z2}, sel, n > \in \texttt{Es/} \land n_{z1} \neq n_{z2}
   \]
   \texttt{is\_sel/} becomes “true” for node “\texttt{n}” and selector “\texttt{sel}” when there are two different nodes \texttt{n_{z1}} and \texttt{n_{z2}} which are compatible (using the \texttt{compatible/}) with “\texttt{n}” and both of them reference the node “\texttt{n}” by selector “\texttt{sel}”.
   We extend the semantic of this statement as follows:
   \[
   \texttt{is\_sel/}(n, \texttt{sel}) =
   \begin{align*}
   &\texttt{is\_sel/}(n, \texttt{sel}) \lor \texttt{is\_sel/}[\texttt{Es/}](n, \texttt{sel}) \\
   &\quad \text{if } \exists n_{x} \in \texttt{Ev/} \land n_{x} > \in \texttt{Ev/} \text{ and } n_{x} \in \texttt{Ev/} \\
   &\quad \text{if } [y, n] \in \texttt{Ev/} \\
   &\quad \text{otherwise}
   \end{align*}
   \]

That is, after breaking references to nodes pointed to by variable “\texttt{x}” using selector “\texttt{sel}”, we check whether or not these referenced nodes maintain the shared attribute for selector “\texttt{sel}”.

5. Statement \([x, \texttt{sel} = y] \)
   This statement can change the \texttt{is\_sel/} information of the nodes directly pointed to by variable “\texttt{y}”, since they are going to be referenced by selector “\texttt{sel}” from the heap.
   \[
   \texttt{is\_sel/}(n, \texttt{sel}) =
   \begin{align*}
   &\texttt{is\_sel/}(n, \texttt{sel}) \lor \texttt{is\_sel/}[\texttt{Es/}](n, \texttt{sel}) \quad \text{if } [y, n] \in \\
   &\quad \text{if } \texttt{Ev/} \\
   &\quad \text{otherwise}
   \end{align*}
   \]

With the \texttt{is\_sel/} function we check if the nodes pointed to by variable “\texttt{y}” are referenced more than once by selector “\texttt{sel}”. The others nodes do not change.

6. Statement \([x := y\texttt{sel}] \)
   The changes induced by this statement are twofold. First, we note that the \texttt{is\_sel/} attribute does not change. However, like in the statement \([x := y] \), we need to take into account the new nodes, which are now pointed to by variable “\texttt{x}”.
   \[
   \texttt{is\_sel/}(n_{x}, \texttt{sel}) = \texttt{is\_sel/}(n_{\texttt{null}}(x), \texttt{sel}) \forall \texttt{seli}
   \]
On the other hand, the `is_set` attribute must be taken into account during the node materialization in order to avoid the creation of unnecessary references. Therefore, we have modified the following functions:

- **compatible**([y, ny], < ny, sel, n_z >):

  compatible((ny, nz, nw) \& \& (y, ny) \in Es \& \&
  < ny, sel, nz >, < ny, sel', nz > \in Es \& \&
  (ny = # \& \& sel = sel') \lor is_set(n_z, sel))

  Note that in this previous function, we use `is_set` instead of `is`#. This function, `compatible`, is used to set new references from the already existing nodes to the materialized one. These new references are of two types: (a) references from selector “sel” from nodes pointed to by “y” variable (corresponding to the statement [x := y; sel]), and (b) other node references, which are going to be taken into account only if `is_set(n_z, sel)` is true.

- **compatible**([y, ny], < ny, sel, n_z >):

  compatible((ny, nz, nw) \& \& (y, ny) \in Es \& \&
  < ny, sel, nz >, < ny, sel', nz > \in Es \& \&
  (ny = # \& \& sel = sel') \lor is_set(n_z, sel))

  Similarly to `compatible` function, we use `is_set` instead of `is`. This function creates “self references” in the materialized node. In order to do this, we only consider those references for which node n_z is shared for selector “sel”.

### 4.4 Cycle links

In order to reduce the number of unnecessary edges in the SSG, we assign a new attribute to each node: `cycles`. This attribute is actually a set of pairs of references < sel1, sel2 >. For a certain node, the pairs in `cycles` fulfill the following property: when taking sel1 and sel2 subsequently from this node, the resulting reference points to the original node. This set maintains similar information to that of “identity paths” in the Abstract Storage Graph (ASG) [11], which is very useful for dealing with doubly linked structures.

Again, the following modifications of the abstract interpretation are needed:

1. Statement [x := nil]

   This statement may produce the summarization of the nodes pointed to by variable “x”. When two nodes are joined, we keep the compatible `cycles` of both of them. A “cycle link” < sel1, sel2 > belonging to `cycles` (n1), is compatible with `cycles` (n2), if (1) < sel1, sel2 > belongs to `cycles` (n2) as well, or (2) the node “n2” does not reference any node by selector “sel1”.

   `cycles` (n1) = { < sel1, sel2 > |< sel1, sel2 >\in
   (cycles (n2), cycles (n2 \cup \{x\})) \lor
   < sel1, sel2 >\in cycles (n2) \land
   \neg \exists n, < n, sel1 >, n \in Es \lor
   < sel1, sel2 >\in cycles (n2) \land
   \neg \exists n, < n, sel1 >, n \in Es \}

2. Statement [x := new]

   For this sentence we create a new node with an empty `cycles`.

   `cycles` (n0) = ∅

3. Statement [x := y]

   The `cycles` of the nodes pointed to by “y” are preserved. In addition, these nodes are also pointed to by “x”.

   `cycles` (n1) = cycles (n0)

4. Statement [x; sel := nil]

   This statement results in the deletion of some elements in the `cycles` set. First, elements of type < sel, sel > of `cycles` (n1) are deleted, where n1 refers to the nodes directly pointed to by the “x” variable. On the other hand, we also delete elements < sel1, sel > from the `cycles` (n0), where n0 are the nodes referenced from n1 by selector “sel”. For this case, n0 should reference n1 by “sel1”.

   `cycles` (n1) =
   - `cycles` (n1) \& < sel, sel1 >
     if < x, n > \in Es \& \& < sel, sel1 > > cycles (n1)
   - `cycles` (n1) \& < sel, sel >
     if < x, n > \in Es \& \& < n, sel1 > > Es \& \&
     < sel, sel1 > > cycles (n1)
   - `cycles` (n1) otherwise

---

Figure 6: (a) Doubly linked list. (b) Shape Graph without `is_set` attributes. (c) Shape Graph with `is_set` attributes. In (c), when a loop traverses the List only by selector “nxt” or “prv”, can we conclude that the same node cannot be visited twice.
5. Statement $[x \text{.sel} = y]$

This statement may create elements in the $\text{cyclelinks}^\#(n_X)$ and $\text{cyclelinks}^\#(n_Y)$ sets. Here, $n_X$ is pointed to by "x" and $n_Y$ is pointed to by "y". Regarding $\text{cyclelinks}^\#(n_X)$, we extend this set with <$\text{sel}, \text{seli}$> elements if $n_Y$ only points, by selector "sel", to nodes directly pointed to by variable "x". In a similar way, we include <$\text{sel}, \text{sel}$> in $\text{cyclelinks}^\#(n_Y)$.

$\text{cyclelinks}^\#(n) =$

- $\text{cyclelinks}^\#(n) \cup <$<$\text{sel}, \text{seli}$>
  if $[x, y], [y, n] \in \text{Ev}^\# \wedge \text{compatible}(n, n_Y) \wedge$
  $<n, \text{sel}, n > \in \text{Es}^\# \land$
  $\neg \exists n_Z \in \text{cyclelinks}^\#(n, n_Z), n \neq n_Z, <n, \text{sel}, n > \in \text{Es}^\#$

- $\text{cyclelinks}^\#(n) \cup <$<$\text{sel}, \text{sel}$>
  if $[x, n], [y, n] \in \text{Ev}^\# \wedge \text{compatible}(n, n_X) \wedge$
  $<n, \text{sel}, n > \in \text{Es}^\# \land$
  $\neg \exists n_Z \in \text{cyclelinks}^\#(n, n_Z), n \neq n_Z, <n, \text{sel}, n > \in \text{Es}^\#$

- $\text{cyclelinks}^\#(n)$ otherwise

6. Statement $[x := y.\text{sel}]$

The $\text{cyclelinks}^\#$ of the nodes pointed to by "y.\text{sel}" are preserved. In addition, these nodes are also pointed to by "x". The materialized nodes will have the same set of "cycle links" as the node from which it has been materialized.

$\text{cyclelinks}^\#(n_Z) = \text{cyclelinks}^\#(n_{Z-\{x\}})$

Once we have applied all these updates in the SSG, it is necessary to check whether or not the references in the graph correspond to the information provided by the cycle links sets. Actually, the method breaks the references which are not compliant with the cycle links sets.

Let be

$A = \{ n | [(x, n) \in \text{Ev}^\#] \lor (<n_x, \text{sel}, n > \in \text{Es}^\#) \lor (<n, \text{sel}, n_x > \in \text{Es}^\#), [x, n] \in \text{Ev}^\#]\}$

The new set of selector edges is

$\text{Es}^\# = \text{Es}^\# \setminus \{<n, \text{sel1}, n_Z > | n \in A, <n, \text{sel1}, n_Z > \in \text{cyclelinks}^\#(n),<n, \text{sel1}, n_Z > \in \text{Es}^\#, <n, \text{sel2}, n > \notin \text{Es}^\#\}$

In Fig. 7 we can see an example showing the improvement that can be achieved by the use of cycle links set. In Fig. 7 (a) we show a doubly linked list with two nodes and their corresponding cycle links sets. Figures 7 (c) and (b) show the resulting SSG after executing the sentence "x:= list.next" , taking the cycle link information into account or not, respectively. We can see that there are two artificial references in case (b), that can be avoided by considering the cycle links information (c), which leads to more accurate SSG's.

4.5 Sparse LU SSG modified

All these previously described techniques have been implemented in a simple compiler which reads C code and returns the SSG for each program point. The compiler has been written in C, taking special care over memory management and in the selection of a proper data structure to store the SSG.

Our Sparse LU factorization in C is transformed to fulfill the normalization assumptions according to the abstract semantic of the SSG method:

- Only one constructor or selector is applied per assignment statement.
- All allocation statements are of the form $x := \text{new} (x, \text{sel} := \text{new} \text{ is not allowed})$.
- In each assignment statement, the same variable does not occur on both the left-hand and right-hand side.
- Each assignment statement of the form $\text{lhs} := \text{rhs}$ in which $\text{rhs} \neq \text{nil}$ is immediately preceded by an assignment statement of the form $\text{lhs} := \text{nil}$.

The equivalent statements in C for the six kinds of statement presented before are: $x = \text{NULL}, x := \text{sel} = \text{NULL}, x = \text{allocate}(), x = y, x := \text{sel} = y$ and $x = y := \text{sel}$.

The resulting SSG for this code is shown in Fig. 8. As we can see, variable A points to a doubly linked list, summary node $n_{k1}$, with the shared attribute set to false. Every node in this list points to a different doubly linked list, represented by summary node $n_{k2}$, by selector "head". There are two summary nodes, because there are two types of pointers ("head list" and "element list").

For all nodes, $i_s$.$\text{sel}^\#(n, \text{sel})$ is "FALSE" for all selectors. Therefore, we conclude that the "head list" and the "element list" are acyclic structures when they are traversed by a single selector type. In addition, we note that there is no shared node among different "element lists". With all these results, it is clear that several sparse matrix columns can be...
updated in parallel during factorization. Furthermore, the resulting SSG points out that each column (element list) is acyclic when traversed using a single selector type. So, if they are actually traversed in this way, it is also possible to update each column in parallel.

For instance, our sparse data structure is traversed in our algorithm by using the "nxt" selector. The "prv" selector is only used to simplify the deletion operation of an entry. In other words, "nxt" can be seen as a "traversing link" whereas "prv" as a "referencing link". However, this "traversing" information should be inferred in a subsequent compiler stage which uses the SSG to perform the data dependence test step. We do not cover this step yet, but we will address this topic in the near future.

It is important to note here that the same SSG we see in Fig. 8 is achieved in two program points: after the data structure initialization (initial sparse matrix A) and after the sparse LU factorization (data structure for the factorized matrix LU). In other words, the in-place LU factorization code does not change the sparse data structure representation and characteristics.

Compared to the SSG method presented by Sagiv [12], we see that our SSG is more accurate than the one they obtain (already presented in Fig. 3). The main reasons leading to their results are: (a) they can only represent a summary node in the SSG whereas we can include many of them for each structure type, (b) their summary node has is@# = true due to the fact that their method is not able to detect that there is not more than one reference from the heap to each node by the same selector. Furthermore, by considering the "is@ed" and "cycle_links" attributes, we are able to infer that there are no shared elements in the list, and also we keep the SSG as accurate as possible avoiding superfluous references.

5 Conclusions and future work

In this work, we have implemented a Shape-Analysis algorithm based on the method of Sagiv et al. [12] (SSG), and that of Plevyak et al. [11] (ASG), improving the accuracy and dealing with more complex data structures.

We have validated the implementation of the method with a real code, the Sparse LU Factorization, for which we have achieved good results. The resulting shape graph provides a great deal of information at compile time. Summarizing, this information describes the data structure used in the algorithm, stating that the columns of the sparse matrix are stored in memory as doubly linked lists which do not share elements. A subsequent data dependence test phase would determine that the algorithm traverses the columns of the reduced submatrix for each k-loop iteration (see Fig. 1) and that each column can be updated in parallel.

This data dependence phase is one of the topics on which we need to focus next. But first, we plan to enhance the method in order to handle more complex data structures, like Acyclic Direct Graphs (ADG) where more than one reference to a node by the same selector may exist. In addition, we are working to extend the shape analysis algorithm to make an interprocedural analysis without applying inlining. Since there are many codes which use recursive calls to traverse the data structure, this is also an important topic.

References