

1 Pointer analysis as a graph problem

Here, we will try to model pointer analysis as a graph theoretic problem. We will try to see if we can get a different formulation from the existing analyses and try to do some optimisations.

Every graph is defined by a set of vertices and edges. We need to define a particular set of vertices and edges to formulate the points to analysis as a graph problem.

1.1 A natural attempt

We can use the graph defined by the points-to relation. Suppose $G = (V, E)$ is a directed graph. Then,

V = The set of variables (pointers and pointees)

$E = \{(p, q) \mid p \text{ points to } q\}$

Some observations

- If our language is strongly typed, then G will be a DAG. Furthermore, edges will only go from a vertex of type (T^*) to (T) (Figure 1).
- We can merge two vertices if they have the same *points-to* and *pointed-by* sets.

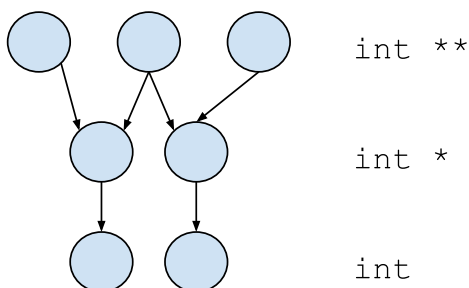


Figure 1: A sample points-to graph for a strictly typed language

1.2 Constraint graph

Consider the following formulation which mimics Andersen's analysis. Suppose $G = (V, E)$ is a directed graph. Then,

$V = \text{The set of pointers}$

$$E = \{(p, q) \mid \text{PtsTo}(p) \subseteq \text{PtsTo}(q)\}$$

where $\text{PtsTo}(p)$ denotes the points to set of p .

We have the following algorithm.

Algorithm 1

Input: Set C of points-to constraints

Output: Points-to information

- 1: Process address-of constraints
 - 2: Add edges to constraint graph G using *copy* statements ¹
 - 3: **repeat**
 - 4: Propagate points-to information in G
 - 5: Add edges to G using load and store statements
 - 6: **until** Fixed point
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Example 1. Consider the following constraints.

$e = d; b = a;$

(Copy constraints)

$*e = c; c = *a; *a = p;$

(Load/store constraints)

$a \mapsto \{a, q, r, s, t\}, p \mapsto \{b, c, d\}$

(Initial points-to sets)

Figure 2 shows each iteration of the above algorithm on this input. The nodes q, r, s, t have been combined into one in the figure for clarity because we know that they have the same points-to sets at the end of the algorithm (*after having computed it once before!*).²

Till now, it seems as if the graph formulation is identical to Andersen's analysis. Then what benefits does such a formulation provide?

- A naive formulation offers no benefits over the constraint-based formulation.
- We need to exploit structural properties of the constraint graph for efficient execution.
 - Online cycle detection
 - Online dominator detection
 - Propagation order: Topological sort, depth first

1.2.1 Pointer Equivalence

Definition 1 (Pointer Equivalence). *Two pointers p and q are equivalent if they have the same points-to set (at the end of the analysis).*

¹Note that we only need to process copy statements once, as opposed to Andersen's analysis

²Interestingly, in this example, we can see that the input is symmetric with respect to q, r, s, t . Suppose we interchange q and r (or any permutation of q, r, s, t), the input as a whole does not change. So one can argue that q, r, s, t will have the same points-to sets without actually running the algorithm.

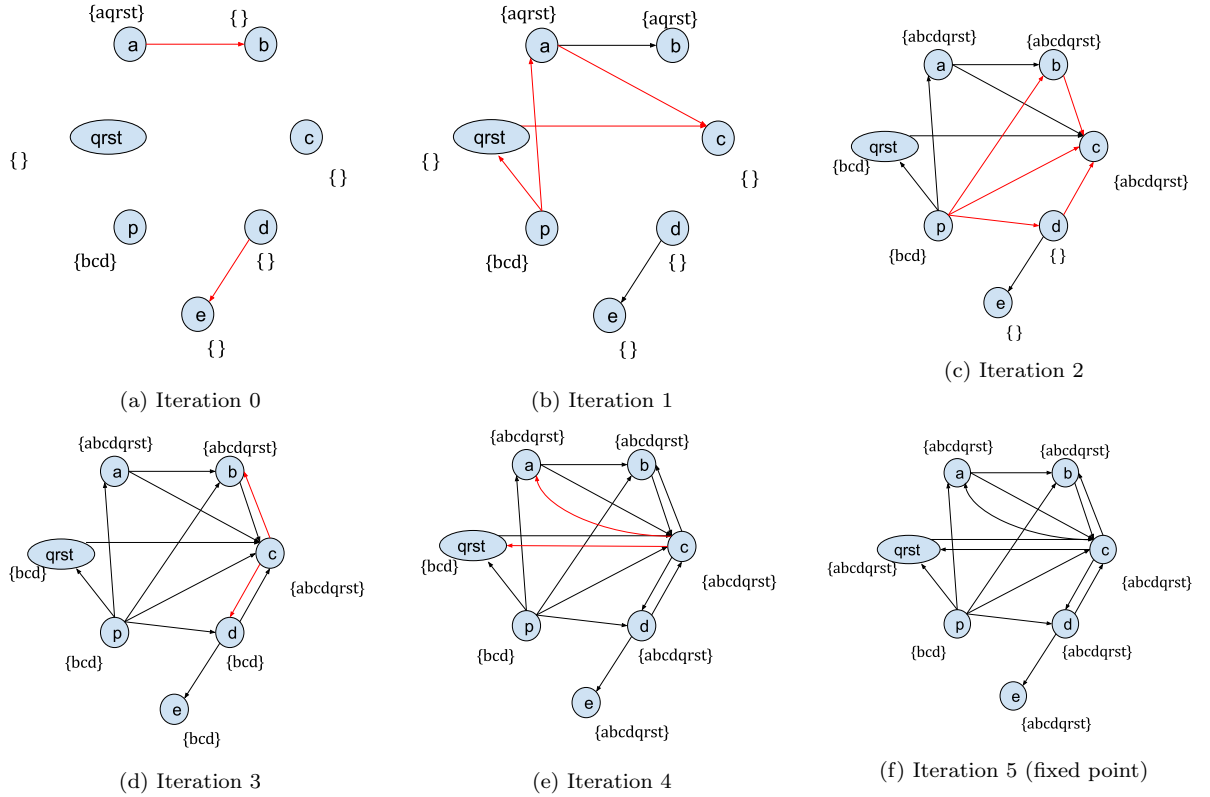


Figure 2: Constraint graph after each iteration as computed by Algorithm 1

Our aim is identify such equivalent pointers *before* computing their points-to information. This will reduce the number of nodes in our constraint graph, and consequently, the amount of information propagated during the analysis, which is usually the costliest operation.

1.2.2 Online cycle detection

Theorem 1. *Suppose the pointers $v_1 \dots v_k$ form a cycle in the constraint graph. Then, these pointers are equivalent.*

Proof. From the definition of the constraint graph, $\text{PtsTo}(v_1) \subseteq \text{PtsTo}(v_2) \subseteq \dots \subseteq \text{PtsTo}(v_k) \subseteq \text{PtsTo}(v_1)$. Thus, $\text{PtsTo}(v_1) = \text{PtsTo}(v_2) = \dots = \text{PtsTo}(v_k)$. \square

Since edges are added to the graph dynamically, cycle detection is performed online so that we can *collapse* cycles (due to theorem 1). When we say that we *collapse* cycles, we actually mean that we merge the PtsTo sets, the incoming edges, and the outgoing edges of the pointers in the cycle and replace the whole cycle with a representative element. Since some pointers might have some extra edges added to them, it might seem that this operation would decrease the precision of our analysis. However, this is not the case since the extra edges are not going to flow any *new* information that wouldn't have already flowed through the old edges in the cycle.

We can use the union find data structure to efficiently perform the cycle collapse. The basic idea is that all the pointers in a particular cycle would be placed in the same set and the whole set would be referred by a particular representative pointer. If we find that two sets are connected by a cycle, or if they have a common pointer, then we will merge the two sets. Figure 3 shows an example of how the union-find data structure is used in collapsing cycles.

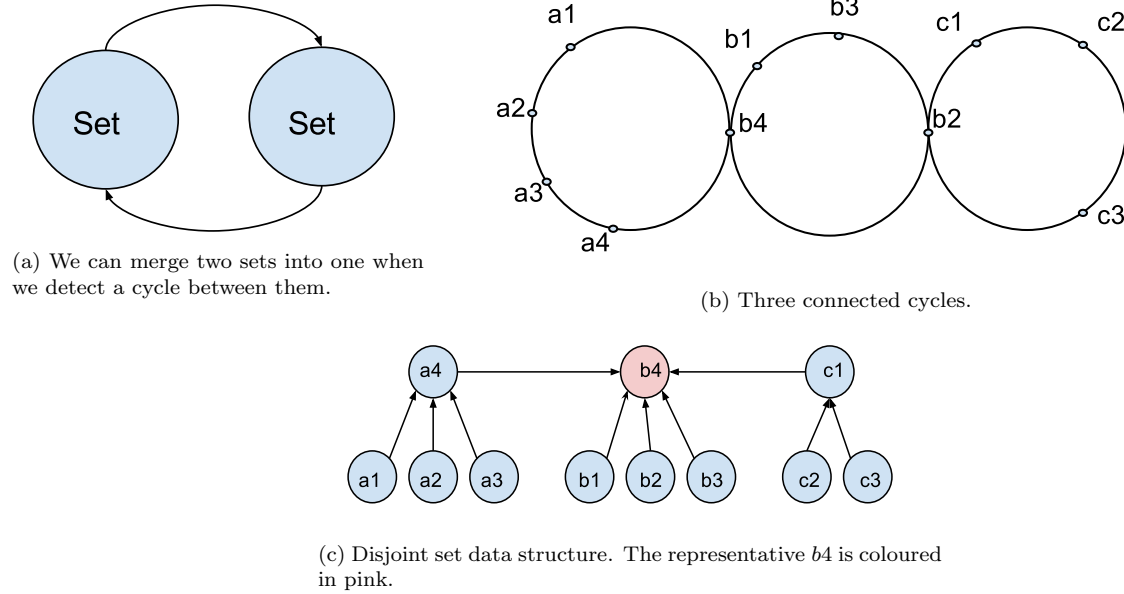


Figure 3: Example showing how disjoint set data structure is used in cycle collapse.

1.2.3 Online dominator detection

Suppose we have a graph structure like that in Figure 4. Let us ignore the address of statements for now. We can see that the only incoming edges to B and C are through the path from A . Then, what can we say about the points-to sets of A , B , and C ? We know that

$$\text{PtsTo}(A) \subseteq \text{PtsTo}(B) \subseteq \text{PtsTo}(C)$$

The only way any information can flow to B and C is through A . So, we can conclude that $\text{PtsTo}(A) = \text{PtsTo}(B) = \text{PtsTo}(C)$.

Note: We can only perform this optimisation if the graph structure at the *end* (fixed point) looks like this. If some other incoming edge gets added to B or C during the analysis and we keep the nodes merged, then our analysis will become imprecise.

We will now formalise this notion of ‘no other incoming edge other than A ’.

Definition 2 (Domination). *In a directed graph with a start node S , we say that a node A dominates B (written as $A \text{ dom } B$) if every path from S to B must go through A .*

Remark: dom is a transitive relation.

Theorem 2. *Suppose we define an appropriate start node for our constraint graph and ignore the address-of constraints.³ Then we have the following:*

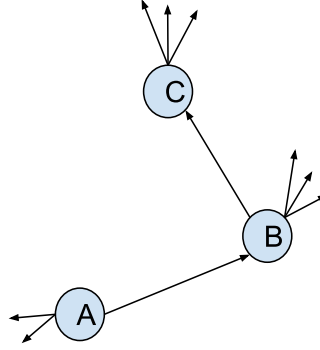


Figure 4: Note that B and C don't have any other incoming edges other than through A

1. If two nodes in a constraint graph have the same dominator, they are pointer equivalent.⁴
2. A dominator and its dominees are pointer equivalent.

Proof. Suppose $A \text{ dom } B$ and $A \text{ dom } C$. We know that $\text{PtsTo}(A) \subseteq \text{PtsTo}(B)$ and $\text{PtsTo}(A) \subseteq \text{PtsTo}(C)$. By definition of **dom**, any information that flows into B or C , must pass through A , so they cannot have any 'extra' information which is not present in A . Therefore, we can say that $\text{PtsTo}(A) = \text{PtsTo}(B)$ and $\text{PtsTo}(A) = \text{PtsTo}(C)$. \square

If we want to use this optimisation to merge some nodes in our graph, we need to be able to perform dominator detection online (during the analysis). Furthermore, we need to make sure that once we merge a set of nodes, some more edges are not added later to the set in such a way that it changes the **dom** relation within the set.

Can we make sure that a particular **dom** relation (say, $A \text{ dom } B$) will never change? In other words, once a dominator, always a dominator. Here is a sufficient condition: if we can ensure that B has only certain fixed predecessors, then we can ensure the above condition. So, in our algorithm, after all copy statements are processed the only way any new incoming edge can be added to B is through a statement of the type $*P = Q$, where B is present in the points-to set of P . Now, if B 's address is never taken, i.e. there is no statement of the type $P = \&B$, then we can say that such a load statement will not add any incoming edge to B .

1.2.4 Offline Variable Substitution

Some constraints are easy to check without running the analysis, such as

- $a = b, b = a$: This forms a cycle between a and b .
- $a = *p, *p = a$: This forms a cycle between a and the elements in the points-to set of p .

³We can model the address-of constraints by adding a node ' $\&x$ ' for every variable whose address has been taken and adding the edge $\&x \rightarrow p$ for every statement of the form $p = \&x$. We can then define a start node S and add edges from S to every address-of node.

⁴This does not hold if we choose the start node S as our dominator, or else, every variable would have the same points-to set.

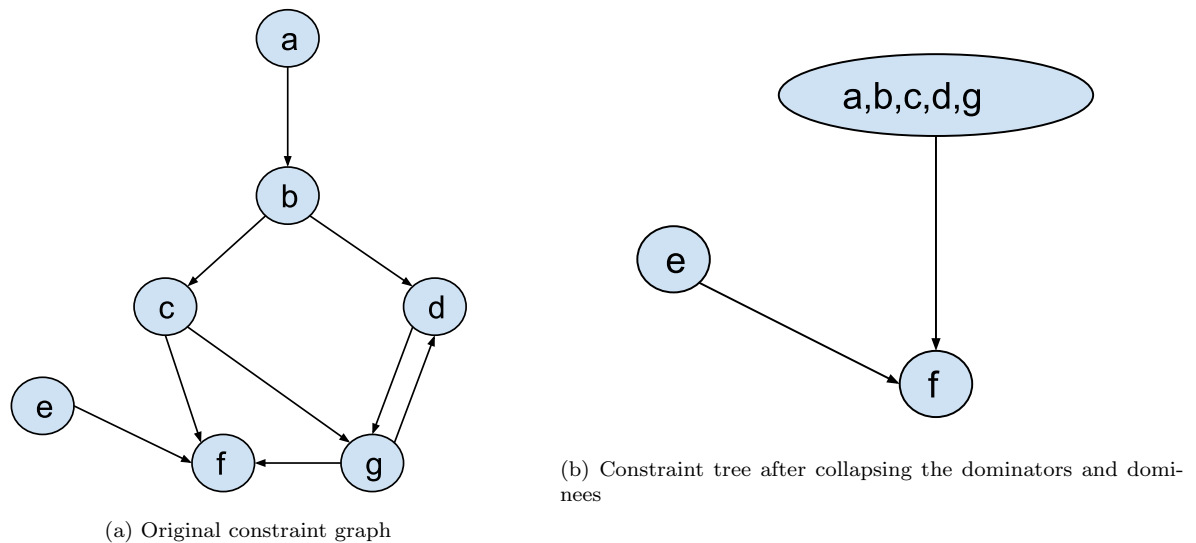


Figure 5: An example showing the dominator-dominee collapse

- $a = b$, $c = a$, $c = b$ and no other incoming edge to c : This creates the $a \text{ dom } c$ relation.

Offline variable substitution is performed before running the pointer analysis. We need to perform OVS in such a way that the total time and space of the analysis (including OVS) is less than the time taken in the analysis without any OVS.

1.2.5 Propagation Order

1. **Wave propagation:** A topological ordering is beneficial for propagating points-to information. It may reduce the time taken per iteration. This is called wave propagation.
2. **Deep propagation:** Wave propagation may consume a lot of memory. In such cases, propagating information in a depth-first manner can reduce the memory requirement. Only the differences may also be propagated to further reduce the memory usage. It also takes advantage of the locality of reference. This is called deep propagation.

References

- [1] R. NASRE, CS6843 course slides
- [2] R. NASRE, Exploiting the structure of the constraint graph for efficient points-to analysis