# Dominators in Graphs

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# Definitions

d dominates n

```
d \operatorname{dom} n, \quad d \in \operatorname{Dom}(n)
```

- If every path from  $s_0$  (start node) to n contains d.
- Every node dominates itself
- d strictly dominates n
  - If d dominates n and  $d \neq n$
- d immediately dominates n d = idom(n)

• If d strictly dominates n and every other dominator of n dominates d

Dominator tree

- Every child is immediately dominated by its parent
- Root node is the start node
- Ancestor a of node  $n \rightarrow a \operatorname{dom} n$



FIGURE 18.3. (a) A flow graph; (b) its dominator tree.

# Applications

- Static Single Assignment
  - Compute the *dominance frontier* first
  - Use the frontier to insert phi nodes
- Finding loops
- Hoisting instructions
- And many other optimizations...

# Anecdotal Performance in Ř

	Before		After	
	Total	Average	Total	Average
Benchmark (5 iterations)	89,205 ms	17,841 ms	82,425 ms	16,485 ms
Constructing the dominator tree	4,988 ms	80 µs	90 ms	1.5 μs
a dom? b	42 ms	46.1 ns	169 ms	182 ns
<i>a</i> idom? <i>b</i>	186 ms	24 ns	153 ms	19 ns

# History

- 1959: Definition of *dominance* [Prosser 1959]
- 1969: First sketch of algorithm [Lowry and Medlock, 1969]
  - Complexity is at least quadratic
- 1970: Data-flow equations [Allen, 1970]
- 1972: Iterative data-flow algorithm [Allen and Cocke, 1972] ... more quadratic algorithms ...
- 1979: Almost linear complexity [Lengauer and Tarjan, 1979] ... more almost linear algorithms ...

### Data-flow Equations

Let Dom(n) be the set of nodes that dominate n. Then:

 $Dom(s_0) = \{s_0\}$ 

$$Dom(n) = \{n\} \cup \left(\bigcap_{p \in preds(n)} Dom(p)\right)$$

Iterate until you reach a fixed point.  $O(n^2)$  time complexity, and very slow in practice

#### Another Algorithm [Aho and Ullman, 1972]

For each node  $v \neq s_0$ :

- Remove v from the graph
- Consider the set of nodes *S* that are now unreachable
- Then Dom(v) = S

 $O(n^2)$  time complexity

### Iterative Algorithm, revisited

- Set intersection is the bottleneck
- Idea: use a consistent ordering for sets
  - Perform intersection by walking through both sets in order, with pairwise comparisons
  - Order encodes a path through the dominator tree
- Very simple implementation [Cooper, Harvey, and Kennedy, 2006]
- But slower in practice than the Lengauer-Tarjan algorithm [Georgiadis, Tarjan, and Werneck, 2006]

# Lengauer-Tarjan Algorithm

- Simple version:  $O(m \log n)$
- Sophisticated version:  $O(m \alpha(m, n))$ 
  - Where  $\alpha(m, n)$  is the inverse of the Ackerman function

The simple Lengauer-Tarjan algorithm is faster in practice, and less sensitive to pathological graphs.

This explanation is adapted from Appel and Palsberg [2004].

# Depth-First Spanning Tree

- Use DFS to compute a spanning tree
  - Assign a *dfnum* to each node

*a* is an *ancestor* of *b* 

- If a = b or there is a path from a to b in the spanning tree
- *i.e.* dfnum(a)  $\leq$  dfnum(b)
- *a* is a *proper ancestor* of *b* 
  - If a is an ancestor of b and  $a \neq b$
  - *i.e.* dfnum(*a*) < dfnum(*b*)

Note: Can test ancestor relation by comparing *dfnum*s



[Appel and Palsberg, 2004]

# Dominators and *dfnum*s

If idom(n) = d, then d must be an ancestor of n *i.e.*, dfnum(d) < dfnum(n)</li>

- Therefore: ancestors of *n* are idom candidates!
- If an ancestor x does not dominate n, there must be some "detour" starting above x.
  - Nodes on the detour are not ancestors of *n*
  - *i.e.* their *dfnums* must be greater than *n*'s



#### Semidominators

s semidominates n

$$s = semi(n)$$

- If *s* is the highest ancestor with a path to *n*, using non-ancestor nodes
  - Highest ancestor → smallest dfnum
    - dfnum(s) < dfnum(n)
  - Path  $p = s, u_1, ..., u_k, n$  using non-ancestor nodes
    - dfnum $(u_i) > dfnum(n)$

#### semi(n) is a candidate for idom(n)

- Often, semi(n) = idom(n)
- An exception: semi(n) itself is bypassed



# Semidominator Theorem

Consider all predecessors v of n in the CFG. Then:

- If v is a proper ancestor of n dfnum(v) < dfnum(n)
  - *v* is a candidate for semi(*n*)
- If v is a non-ancestor of n dfnum(v) > dfnum(n)
  - For each u that is an ancestor of v, (and not an ancestor of n) semi(u) is a candidate for semi(n)

semi(n) is the candidate with the lowest *dfnum* 





#### **Dominator Theorem**

Consider the spanning tree path from s = semi(n) to n. Let y be the node with smallest numbered semidominator, *i.e.* minimum dfnum(semi(y)).

$$idom(n) = \begin{cases} semi(n) & \text{if } semi(y) = semi(n) \\ idom(y) & \text{if } semi(y) \neq semi(n) \end{cases}$$



[Appel and Palsberg, 2004]

# Lengauer-Tarjan Algorithm

- 1. Perform DFS to number nodes and create the depth-first spanning tree
- 2. For each node *n* (in decreasing *dfnum* order):
  - Use the Semidominator Theorem to compute semi(n)
  - Insert *n* into the spanning forest
- 3. Implicitly define the idom by applying the first clause of the Dominator Theorem
- 4. For each node *n* (in increasing *dfnum* order):
  - Explicitly define the idom by applying the second clause of the Dominator Theorem

# Spanning Forest

- Build a spanning forest as the CFG is traversed
  - When *n* is processed, only non-ancestors of *n* are in the forest
- •link(p, n)
  - Add the edge (n, p) to the spanning forest
- ancestorWithLowestSemi(v)
  - Search upwards in the forest, starting from v
  - Find the ancestor of *v* whose semidominator has the lowest *dfnum*





**FIGURE 19.11.** Path compression. (a) Ancestor links in a spanning tree; AncestorWithLowestSemi(v) traverses three links. (b) New nodes  $a_2$ ,  $a_3$  are linked into the tree. Now AncestorWithLowestSemi(w) would traverse 6 links. (c) AncestorWithLowestSemi(v) with path compression redirects ancestor links, but *best*[v] remembers the best intervening node on the compressed path between v and  $a_1$ . (d) Now, after  $a_2$  and  $a_3$  are linked, AncestorWithLowestSemi(w) traverses only 4 links.

```
Dominators() =
   N \leftarrow 0; \forall n. bucket[n] \leftarrow \{\}
   \forall n. dfnum[n] \leftarrow 0, semi[n] \leftarrow ancestor[n] \leftarrow idom[n] \leftarrow samedom[n] \leftarrow none
   DFS(none, r)
   for i \leftarrow N - 1 downto 1
        n \leftarrow vertex[i]; p \leftarrow parent[n]; s \leftarrow p
        for each predecessor v of n
             if dfnum[v] \leq dfnum[n]
                s' \leftarrow v
             else s' \leftarrow semi[AncestorWithLowestSemi(v)]
             if dfnum[s'] < dfnum[s]
                s \leftarrow s'
        semi[n] \leftarrow s
        bucket[s] \leftarrow bucket[s] \cup \{n\}
        Link(p, n)
        for each v in bucket[p]
             y \leftarrow \text{AncestorWithLowestSemi}(v)
             if semi[y] = semi[v]
                idom[v] \leftarrow p
             else samedom[v] \leftarrow y
        bucket[p] \leftarrow \{\}
   for i \leftarrow 1 to N-1
        n \leftarrow vertex[i]
        if samedom[n] \neq none
           idom[n] \leftarrow idom[samedom[n]]
```

```
DFS(node p, node n) =

if dfnum[n] = 0

dfnum[n] \leftarrow N; \quad vertex[N] \leftarrow n; \quad parent[n] \leftarrow p

N \leftarrow N + 1

for each successor w of n

DFS(n, w)
```

AncestorWithLowestSemi(node v) =  $a \leftarrow ancestor[v]$ if  $ancestor[a] \neq$  none  $b \leftarrow$  AncestorWithLowestSemi(a)  $ancestor[v] \leftarrow ancestor[a]$ if dfnum[semi[b]] < dfnum[semi[best[v]]]  $best[v] \leftarrow b$ return best[v]Link(node p, node n) =  $ancestor[n] \leftarrow p; \quad best[n] \leftarrow n$ 

[Appel and Palsberg, 2004]

# Implementation in Ř

Straightforward translation from pseudocode to C++

• About 100 LOC, without comments

https://github.com/reactorlabs/rir/blob/b265f9e/rir/src/com piler/util/cfg.cpp#L52