Jive Research Overview Towards Scalable Visualizations

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Oiagram Filtering



- Oiagram Filtering
- 4 Lifeline Projection



- 2 Diagram Folding
- Oiagram Filtering
- 4 Lifeline Projection
- Combining Techniques



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- Conclusion and Future Work



Diagram Folding

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A Closer Look at Sequence Diagrams

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 - Compose objects to create more complex interactions.
 - Instantiate large number of objects to accomplish a program's tasks.
 - Have long execution histories- even programs of modest size!

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 - Grow vertically (complex iterations means more method activations).

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- As a result, sequence diagrams tend to:
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- Bottom line: sequence diagrams quickly become unmanageable.

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 - Filter out predefined sets of calls and returns.
 - Project the sequence diagram along specified lifelines.
 - Combine one or more of the above.

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- For all of the above, we must define appropriate criteria!



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Diagram Folding: Agenda

- Define an adequate data structure to abstract the sequence diagrams.
 - Our key data structure is the Call Tree.
 - We maintain one call tree per execution thread.
- Identify useful folding criteria.
 - Along the sequence diagram depth (fold subtrees).
 - Along the sequence diagram breadth (fold adjacent siblings).
- Define the necessary folding operations.

Call Tree

- Directed tree.
- Nodes correspond to method activations and edges to method calls.
- Every node *n* has an associated tuple $\tau(n) = \langle m, e, c, r \rangle$, where:
 - *m* is the called method,
 - e is the method's execution environment (e.g., an object or a class),
 - $c \in \mathbb{N}$ is the method's call time, and
 - $r \in \mathbb{N} \cup \{\pi\}$ is the method's return time.
- Edge (n_1, n_2) encodes a method call from n_1 (caller) to n_2 (callee).
- Total order '<' on call tree nodes: $n_1 < n_2 \Leftrightarrow \tau(n_1).c < \tau(n_2).c$.
- Observation: one call tree per thread!











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Operations

• Fold(t,f)

- replaces the subtree rooted at f in t with a new leaf node ℓ .
- FoldBefore(t,n)
 - folds all nodes f of t such that f < n and $f \notin ancestors(n)$.
- FoldAfter(t,n)
 - folds all nodes f of t such that f > n and f \not descendants(n).
- FoldBetween(t,n₁,n₂)
 - folds all nodes f of t such that $n_1 > f > n_2$ and $f \notin descendants(n_1) \cup ancestors(n_2)$.
- Note: call trees traversed breadth first.

Figure: Possible Uninteresting Regions, Assuming Interest in Activations 35 and 76 (dup).





Operations

- In the last example, nodes 5 and 11 could be further folded into a single node.
- One may argue that this is not necessary.
- Yes, for this example there is no obvious benefit.
- What if the folded call tree had 1000 nodes occurring before node 35?
- The FoldXXX procedures presented earlier do not solve this problem.
- We need breadth-wise folding!





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- Regular expressions to the rescue!
- Given any reasonably sized sequence of calls made from the same caller, we assume that there are recurring patterns in the call sequence.
- Recurring patterns are usually due to loops, but may occur due to explicit calling patterns in the body of the caller.

Figure: Diagram Folding- Report.java

```
public class Report {
 1
 2
3
     public String format(String line) { ... }
 4
 5
      public void print(BufferedReader in, PrintStream out) {
 6
 7
       // let N represent the number of lines in the reader
8
       while (String line = in.readLine() != null) {
9
        out.printLn(format(line));
10
11
12
   |}
```

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```

- The sequence of calls in print may be folded to a single node in the call tree.
- The node is labeled with the regex: $(readLine; format; printLn)^{\mathbb{N}}$.
- Such regex compactly represents the call sequence with no loss of information!

- RegexFoldBefore(p,n)
 - computes an ordered list *C* of all child nodes of *p* such that $c \in C \Leftrightarrow c < n$;
 - replaces all children of p occurring in C with a single leaf node ℓ labeled with a regular expression computed from the ordered list of method calls obtained from C.
- RegexFoldAfter(p,n) and RegexFoldBetween(p,n₁,n₂) are defined analogously, with the proper changes to the inclusion criterion of nodes in *C*.
- Note: algorithm for the conversion of the sequence of method calls into a regular expression is not obvious.

- Let Regex(S) be the algorithm that takes a string S as input and returns a regular expression R such that S ∈ R (i.e., S is a string in the language R).
- R is not any regular expression:
 - wildcards and disjunctions are not allowed;
 - only primitive string repeats are allowed: $(ab)^2$ is fine but $(aa)^2$ is not;
 - R is the most compact regex satisfying the above criteria (need not be unique).
- Algorithm:
 - Part I computes all primitive repeats of S in O(|S|·log|S|) time and O(|S|) space;
 - Part II uses Dijkstra's algorithm to compute an optimal regex for S in $O(|S| \cdot log|S|)$ time and space;
 - A simple post-processing step normalizes singletons of R in O(|S|) time and space.





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Filtering

- Filtering is the process by which debug events are omitted from Jive's model.
- As a consequence, call trees no longer contain complete call/return information.
- We introduce **out-of-model** calls and returns to cope with missing information.
- Calls and returns originating and terminating out-of-model are ignored.
- Out-of-model calls to in-model methods are handled as follows:
 - Let n_1 be the largest outstanding in-model node and n_2 the new in-model node.
 - If n₁ has an outstanding out-of-model child o, add n₂ as a child of o;
 - Else, create a new out-of-model node *o*, add *o* as a child of *n*₁, and *n*₂ as a child of *o*.
- All other calls and returns are handled as if no filtering was in place.
- Inferring out-of-model calls and returns is done using call trees and call stacks.
- All algorithms introduced thus far may be adapted to handle out-of-model nodes.
- Sequence diagrams must be extended to handle out-of-model calls and returns.

Figure: Extended Sequence Diagram Arrows. Columns indicate whether calls and returns originate/terminate in-model or out-of-model. The 'in/out+/in' column indicates a call or return originating in-model and terminating out-of-model, followed by any number of out-of-model activations, terminating in-model after a call or return from out-of-model.



Figure: Extended Sequence Diagram for Hanoi Towers. The towers implement toString(), which are called from the out-of-model PrintWriter.format(String, Object, ...).



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Filtering (cont.)

- Jive supports all filtering supported by JPDA:
 - class and package filtering based on regular expressions.
- Jive also implements a number of local filters:
 - method and class filtering based on visibility;
 - method filtering based on regular expressions;
 - synthetic method filtering.
- Jive provides default filters for standard Java and Sun packages.

Lifeline Projection /



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Lifeline Projection /

Lifeline Interactions

- Users may be interested in only a subset of lifelines in the sequence diagram.
- Users may project any number of selected lifelines into a new sequence diagram.
- The new diagram only contains method activations occurring in these lifelines.
- Calls and returns from other lifelines are represented as out-of-model.
- Lifelines may be reordered during projection to minimize edge crossings.

Combining Techniques /



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Combining Techniques /

Visualizing Query Results

- Query results are typically represented as points in the sequence diagram.
- The FoldXXX and RegexFoldXXX procedures reduce the diagram's depth and breadth.
- Projecting the lifelines containing result points further reduces the diagram along the relevant objects' lifelines.
- Composing these operations automatically yields the smallest relevant sequence diagram for a given query result set.
- Note: these operations may also be combined manually, by the user, and applied to other use cases.

Conclusion and Future Work /



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Conclusion and Future Work /

Status Report

- Currently, Jive supports manual folding of a single node.
- It also fully supports out-of-model calls.
- However, just JPDA filters are currently supported.
- We are in the process of implementing all other algorithms and operations described in here.