Optimized Live Heap Bound Analysis

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Abstract

This paper describes a general approach for optimized live heap space and live heap space-bound analyses for garbage-collected languages. The approach is based on program analysis and transformations and is fully automatic. In our experience, the space-bound analysis generally produces accurate (tight) upper bounds in the presence of partially known input structures. The optimization drastically improve the analysis efficiency. The analyses have been implemented and experimental results confirm their accuracy and efficiency.

1 Introduction

Time and space analysis of computer programs is important for virtually all computer applications, especially in embedded systems, real-time systems, and interactive systems. In particular, space analysis is becoming important due to the increasing uses of high-level languages with garbage collection, such as Java [26], the importance of cache memories in performance [32], and the stringent space requirements in the growing area of embedded applications [29]. For example, space analysis can determine exact memory needs of embedded applications; it can help determine patterns of space usage and thus help analyze cache misses or page faults; and it can determine memory allocation and garbage collection behavior.

Space analysis is also important for accurate prediction of running time [12]. For example, analysis of worst-case execution time in real-time systems often uses loop bounds or recursion depths [25, 2] both of which are commonly determined by the size of the data being processed. Also, memory allocation and garbage collection, as well as cache misses and page faults, contribute directly to the running time. This is increasingly significant as the processor speed increases, leaving memory access as the performance bottleneck.

Much work on space analysis has been done in algorithm complexity analysis and systems. The former is in terms of asymptotic space complexity in closed forms [19]. The latter is mostly in the form of tracing memory behavior or analyzing cache effects at the machine level [24, 32]. What has been lacking is analysis of space usage for high-level languages, in particular, automatic and accurate techniques for live heap space analysis for languages with garbage collection, such as Java, ML or Scheme.

This paper describes a general approach for automatic accurate analysis of live heap space based on program analysis and transformations. The analysis determines the maximum size of the live data on the heap during execution. This is the minimum amount of heap space needed to run the program even if garbage collection is performed whenever garbage is created. This metric is useful for evaluating other garbage collection schemes, just like the performance of an optimal cache replacement algorithm is useful for evaluating other replacement algorithms. The analysis can easily be modified to determine related metrics, such as space usage when garbage collection is performed only at fixed points in the program. It can also be used to help analyze the space usage of some continuously running processes that have cyclic behavior.

Our approach starts with a given program written in a high-level functional language with garbage collection. We construct (i) a space function that takes the same input as the original program and returns the amount of space used and (ii) a space-bound function that takes as input a characterization of a set of inputs of the original program and returns an upper bound on the space used by the original program on

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Form Approved OMB No. 0704-0188 any input in that set. A key problem is how to characterize the input data and exploit this information in the analysis.

In traditional complexity analysis, inputs are characterized by their size. Accommodating this requires manual or semi-automatic transformation of the time or space function [21, 33]. The analysis is mainly asymptotic. A theoretically challenging problem that arises in this approach is optimizing the time-bound or space-bound function to a closed form in terms of the input size [21, 28, 8]. Although much progress has been made in this area, closed forms are known only for subclasses of functions. Thus, such optimization can not be automatically done for analyzing general programs.

Rosendahl proposed characterizing inputs using partially known input structures [28]. For example, instead of replacing an input list l with its length n, we simply use as input a list of n unknown elements. A special value uk ("unknown") is introduced for this purpose. It represents unknown primitive values; if it represented constructed data, we wouldn't know how much space it used. At control points where decisions depend on unknown values, the maximum space usage of all branches is computed. Rosendahl concentrated on proving the correctness of this transformation for time-bound analysis. He relied on optimizations to obtain closed forms, but closed forms can not be obtained for all bound functions.

Our analysis and transformations are performed at source level. This allows implementations to be independent of compilers and underlying systems and allows analysis results to be understood at source level. Our space bound analysis is an abstract interpretation, expressed conveniently as a program transformation. Profiling, like space functions, measures the program's behavior on one input at a time; space-bound functions can efficiently analyze the program's behavior on a set of inputs at once. They can thus be used to determine worst-case space usage of a program, given a particular metric such as input size. Alternatively, worst-case space usage may be determined by applying the space function to a worst-case input. But in general, it is non-trivial to determine such an input. Finding space bounds is undecidable, so space-bound functions may diverge. In our experience, this is rare.

While our approach can be applied to imperative languages, the analysis in this paper enjoys multiple benefits due to the functional nature of the source language. Associating reference counts with partially known structures forms an accurate basis for determining liveness. With reference counting, the heap never has to be examined in its entirety. In contrast, garbage collection algorithms for imperative languages generally examine the whole heap at once which is costly. Space-bound functions sometimes have to evaluate both branches of conditionals due to unknown values. Our analysis does not need to maintain multiple copies of the heap while evaluating the two branches exactly because of the absence of imperative update. Copying the heap would add significant complexity and overhead. It is necessary to keep the results of both branches. This could lead to loose bounds in a naive analysis, since both results seem live. Our analysis handles this by examining selected parts of the heap at a limited number of program points. If the source language were imperative, the analysis would need to examine much larger parts of the heap and do so more frequently.

2 Language

We use a first-order, call-by-value functional language that has literal values of primitive types (e.g., Boolean and integer constants), structured data, operations on primitive types (e.g., Boolean and arithmetic operations), testers, selectors, conditionals, bindings, and function calls. These are fundamental program constructs that have analogues in all programming languages. A program is a set of mutually recursive function definitions of the form $f(v_1, ..., v_n) = e$, where an expression e is given by the grammar

e ::= v	variable reference
l	literal
$ c(e_1,,e_n)$	constructor application
$ p(e_1,,e_n)$	operation on primitive types
c?(e)	tester application
$ c^{-i}(e) $	selector application
if e_1 then e_2 else e_3	conditional expression
$ \mathbf{let} \ v = e_1 \ \mathbf{in} \ e_2$	binding expression
$ f(e_1,, e_n)$	function application

We sometimes use infix notation for primitive operations.

For brevity, we assume the language contains only two kinds of primitive operations that take data constructions as arguments. A tester application c?(v) returns true iff v has outermost constructor c. A selector application $c^{-i}(v)$ returns the i'th component of a data construction v with outermost constructor c. Our analysis can easily be extended to handle other similar operations such as equality predicates.

Input programs to our analysis are assumed to be purely functional, but transformed programs use arrays and imperative update. A sequential composition e_1 ; e_2 returns the value of e_2 . In examples, we use a constructor cons with arity 2.

3 Live Heap Space Function

To analyze the live heap space used by a program on a known input, we transform the program into one that performs all the computations of the original program and keeps track of the total amount of live data. Liveness of data is ascertained using reference counts. The reference count for a data construction v is the number of pointers to v. These may be pointers on the stack, created by let bindings or bindings to formal parameters of functions, or pointers on the heap, created by data constructions. Data construction v is live if its reference count is greater than 0 or if it is the result of the expression just evaluated. Note that in the latter situation, v is still accessible to the program even though there are no references to it.

A constructor count vector v has one element $v[i_c]$ corresponding to each data constructor c used in a given program. Let $P[i_c]$ be the size of an instance of c. Let \cdot denote dot product of vectors. The maximum $\max(v_1, v_2)$ of constructor count vectors v_1 and v_2 is v_1 if $v_1 \cdot P \geq v_2 \cdot P$ and is v_2 otherwise.

The transformation \mathcal{L} in Figure 1 produces live heap space functions. It introduces two global variables, live and maxlive, that satisfy: (1) for each constructor c, live $[i_c]$ is the number of live instances of c; (2) maxlive is the maximum value of live so far during execution. The maximum live space used during evaluation of function f is at most $ml \cdot P$ where ml is the value of maxlive after evaluation of the space or bound function for f.

Our implementation of reference counting is based on an abstract data type (ADT) that defines five functions. $new(c(x_1,\ldots,x_n))$ returns a value v representing a new data construction $c(x_1,\ldots,x_n)$, whose reference count is initialized to zero. data(v) returns the data construction $c(x_1,\ldots,x_n)$. rc(v) returns the reference count associated with v. incrc(v) and decrc(v) increment and decrement, respectively, the reference count associated with v. incrc and decrc are no-ops if the argument is a primitive value.

Updating Reference Counts. rc(v), for a data construction v, is incremented when v is bound to a variable or function parameter, or a data construction containing v as a child is created. rc(v) is decremented when the scope of a **let** binding for v ends, a function call with an argument bound to v returns, or a data construction containing v as a child becomes garbage.

Updating live and maxlive. Whenever new data is constructed, live is incremented, and maxlive is recomputed. An auxiliary function gc ("garbage collect") is called whenever data can become garbage. For a data construction v, gc(v) decrements rc(v) and then, if rc(v) is not positive, it decrements the appropriate element of live and calls gc recursively on the children of v. A data construction may become garbage (1) because of a decrement of its reference count or (2) because it is created in the argument of a selector or tester and is lost to the program after the result of the selection or test is obtained. For example, cons(0,1) is garbage after the application of $cons^{-2}$ in $cons(cons^{-2}(cons(0,1)), 2)$; note that its reference count is always 0.

gcExcept(u,v) is called when u should be garbage collected, v should not be garbage collected and v might be a descendant of u. At the end of function calls and let expressions, values bound to parameters and variables should be garbage collected without garbage collecting the result of the function call or the let expression. Similarly, after selector applications, data selected from should be garbage collected without garbage collecting the selected part, even if the reference count of that part becomes 0. Figure 5 contains an example of a live heap space function.

```
f_L(v_1,\ldots,v_n)=\mathcal{L}[e] where e is the body of function f, i.e., f(v_1,\ldots,v_n)=e
\mathcal{L}[v] = v
\mathcal{L}[l] = l
\mathcal{L}[c(e1,\ldots,en)] = live[i_c] + +;  if (P \cdot live > P \cdot maxlive)
                                                   then for c \in Constructors\ maxlive[i_c] := live[i_c];
                               let r_1 = \mathcal{L}[e_1], \dots, r_n = \mathcal{L}[e_n] in incrc(r_1); \dots; incrc(r_n); new(c(r_1, \dots, r_n))
\mathcal{L}\left[p(e_1,\ldots,e_n)\right] = p(\mathcal{L}\left[e_1\right],\ldots,\mathcal{L}\left[e_n\right])
\mathcal{L}[c?(e)] = \mathbf{let} \ x = \mathcal{L}[e] \ \mathbf{in}
                     let r = c?(data(x)) in
                        (if not(isPrim(x)) and rc(x) = 0 then qc(x)); r
\mathcal{L} \left[ c^{-i}(e) \right] = \mathbf{let} \ x = \mathcal{L} [e] \ \mathbf{in}
                         let r = c^{-i}(data(x)) in
                            (if not(isPrim(x)) and rc(x) = 0 then qcExcept(x, r)); r
\mathcal{L}[\text{if } e1 \text{ then } e2 \text{ else } e3] = \text{if } \mathcal{L}[e1] \text{ then } \mathcal{L}[e2] \text{ else } \mathcal{L}[e3]
\mathcal{L}\left[\mathbf{let}\ v=e_1\ \mathbf{in}\ e_2\right]=\mathbf{let}\ v=\mathcal{L}\left[e_1\right]\ \mathbf{in}
                                       let r = \mathcal{L}[e_2] in
                                          gcExcept(v,r); r
\mathcal{L}\left[f(e_1,\ldots,e_n)\right] = \mathbf{let}\ r_1 = \mathcal{L}\left[e_1\right],\ldots,r_n = \mathcal{L}\left[e_n\right]\ \mathbf{in}
                                  incrc(r_1); \ldots; incrc(r_n);
                                  let r = f_L(r_1, \ldots, r_n) in
                                    gcExcept(r_1, r); \ldots; gcExcept(r_n, r); r
qc(v) = \mathbf{if} \ not(isPrim(v))
              then decrc(v);
                        if rc(v) < 0
                        then live[conType(v)] --;
                                  for i = 1..arity(v) \ gc(c^{-i}(data(v)))
qcExcept(u, v) = incrc(v); qc(u); decrc(v)
```

Figure 1: Transformation that produces live heap space functions f_L . isPrim(v) returns true iff v is primitive. conType(v) returns an integer i_c that uniquely identifies the outermost constructor c in data(v). arity(v) returns the arity of the outermost constructor in data(v).

4 Live Heap Space Bound Function

The transformation \mathcal{L}_b in Figures 2-3 produces live heap space-bound functions. We sometimes refer to space-bound functions simply as bound functions. At every point during the execution of $\mathcal{L}_b[f](x)$, the value of live is an upper bound on the possible values of live at the corresponding point in executions of $\mathcal{L}[f](x')$, for all x' in the set represented by x. As before, maxlive contains the maximum value of live so far during execution. The presence of partially known inputs in bounds analysis causes uncertainty. For conditional expressions whose tests evaluate to uk, both branches are evaluated to determine the maximum live heap space usage.

Correctness of live heap bound analysis depends on keeping track of all references and reference counts meticulously. Summarizing the results of two branches into a single partially known structure that represents both results, as is done in timing analysis [22] and stack space and heap allocation analysis [31], does not work for live heap analysis because it would be impossible to keep track of reference counts accurately. So the result of a conditional whose test evaluates to uk is a separate entity, a join-value, that points to both

possible results and has its own reference count. By keeping both results live, we run the risk of obtaining loose bounds, since *live* might include the sizes of both results when only one of them is live. To keep *live* as tight as possible, we examine join-values at appropriate points of execution and manipulate *live* so that it includes the size of only a single largest data structure pointed to by each join-value.

4.1 Abstract Data Types

Two abstract data types (ADTs), the join-value type and the con-value ("constructed-value") type, are used. A join-value represents a set of possible results. Join-values are created by conditional expressions whose tests evaluate to uk and by selectors applied to join-values. Each join-value j has a list branches(j) containing references to con-values and/or join-values. Primitive values, if any, in the set represented by j are not stored in j. Thus, if branches(j) has only one element, j represents a choice between that element and some primitive value. j has an associated constructor count vector exs(j). Parts of the data constructions represented by j may be live regardless of j. Of the other parts, only those occurring in a single largest branch are live in a worst case (i.e., maximal live heap space) execution of the original program. The sum of the other parts that are not in the largest branch is an excess and is stored in exs(j), as discussed in Section 4.3. live does not include exs(j). When j becomes garbage, exs(j) is added to live just before garbage collecting the branches of j.

The con-value type is an extension of the ADT described in Section 3. con-values and join-values have a reference count and a list of join-parents. A join-value j is a join-parent of v if branches(j) references v. Functions re, incre, decre, joinParents, addJoinParent, and delJoinParent apply to both ADTs; the names indicate their meanings. newjoin(b) creates a join-value j with a list b of branches, and with rc(j) initialized to 0, joinParents(j) initialized to nil, and exs(j) initialized to the zero vector, denoted V_0 .

4.2 Conditionals, Selectors and Testers

Consider a conditional expression (if e_1 then e_2 else e_3)[†] whose condition evaluates to uk. Suppose l_1 , l_2 and l_3 are the values of live after evaluating e_1 , e_1 ; e_2 and e_1 ; e_3 , respectively. The value of live at † is set to $\max(l_2, l_3)$. The result r of the conditional expression is computed by $join(r_2, r_3)$, where r_2 and r_3 are the results of e_2 and e_3 , respectively. If r_2 and r_3 are primitive, then r is r_2 if $r_2 = r_3$ and uk otherwise. If r_2 and r_3 are not primitive and are the same, then r is r_2 . Otherwise, r is a join-value, and exs(r) is set to $\min(l_2 - l_1, l_3 - l_1)$. $l_2 - l_1$ and $l_3 - l_1$ are the amounts of new data in r_2 and r_3 , i.e., the amounts of data created by e_2 and e_3 . e_3 and e_3 may contain old data too, i.e., data created before evaluating e_2 and e_3 . Old data are live regardless of r. Between the sets of new data in r_2 and r_3 , only one set is live. We keep the larger set live; the size of the other set is exs(r).

Observe that in the transformation of conditional (if e_1 then e_2 else e_3), we evaluate e_2 and then e_3 , making copies of only live in between. We do not have to copy the entire heap before or after evaluating either expression because the source language does not contain imperative update. Thus, if h_1 is the heap after the evaluation of e_1 , then e_2 and e_3 modify h_1 only by adding new con-values to it. Informally, h_2 , the heap after evaluation of e_2 , is just $(h_1 + r_2)$, where r_2 is the result of e_2 . Similarly, h_3 is $(h_1 + r_3)$, h_3 and r_3 having the expected meanings. In other words, the heap after evaluation of the conditional is $(h_1 + (r_2 \text{ or } r_3))$. The choice between r_2 and r_3 is conveniently represented using a join-value that points to them both.

Selectors and testers return uk if given uk arguments. For join-values with two non-primitive branches, the selector or tester is first applied to the branches and the join of the results is returned. The exs field of a join-value j that is the result of applying a selector to another join-value j' is set to V_0 , because when j is created, j' is live, and exs(j') already takes care of any excess. In fact, computing exs(j) would yield V_0 as the result, because j' references j's children and as a result, none of the descendants of j are contained-in j.

Applying any tester other than null? to join-values with one primitive branch and one non-primitive branch, results in the join of (a) false (the result of applying any tester other than null? to any primitive branch) and (b) the result of applying the tester to the non-primitive branch. Recall that join-values do not save the values of their primitive branches. Applying null? to an unknown primitive branch yields uk, since the set of primitive values contains both a null value nil and non-null elements. So, the result of null? on

```
f_{Lb}(v_1,\ldots,v_n)=\mathcal{L}_b[e]
                                         where e is the body of function f, i.e., f(v_1, \ldots, v_n) = e
\mathcal{L}_b[v] = v
\mathcal{L}_b[l] = l
\mathcal{L}_b[c(e_1,\ldots,e_n)] = live[i_c] + +;  if (P \cdot live > P \cdot maxlive)
                                              then for c \in Constructors\ maxlive[i_c] := live[i_c];
                            let r_1 = \mathcal{L}_b[e_1], \dots, r_n = \mathcal{L}_b[e_n] in incrc(r_1); \dots; incrc(r_n); new(c(r_1, \dots, r_n))
\mathcal{L}_b[p(e1,\ldots,en)] = p_u(\mathcal{L}_b[e1],\ldots,\mathcal{L}_b[en])
p_u(v_1,\ldots,v_n)= if v_1=uk or \cdots or v_n=uk then uk else p(v_1,\ldots,v_n)
\mathcal{L}_b[c?(e)] = \mathbf{let} \ x = \mathcal{L}_b[e] \ \mathbf{in}
                   let r = c?_u(x) in
                      (if not(isPrim(x)) and rc(x) = 0 then gc(x)); r
c?_u(v) = \mathbf{if} \ v = uk \ \mathbf{then} \ uk
              else if isJoin(v) then if length(branches(v)) = 1
                                                 then join(false, c?_u(first(branches(v))))
                                                 else join(c?_u(first(branches(v))), c?_u(second(branches(v))))
                     else c?(data(v))
\mathcal{L}_b \left[ c^{-i}(e) \right] = \mathbf{let} \ x = \mathcal{L}_b \left[ e \right] \ \mathbf{in}
                       let r = c_u^{-i}(x) in
                          (if not(isPrim(x)) and rc(x) = 0 then gcExcept(x,r); recomputeExs(r)); r
c_u^{-i}(v) = \mathbf{if} \ v = uk \ \mathbf{then} \ uk
             else if isJoin(v)
                     then if length(branches(v)) = 1 then c^{-i}(false)
                              else join(c_u^{-i}(first(branches(v))), c_u^{-i}(second(branches(v))))
                     else c^{-i}(data(v))
\mathcal{L}_b [if e_1 then e_2 else e_3] =
  let b = \mathcal{L}_b[e_1] in
    if b = uk then let l_1 = copy(live) in
                              let r_2 = \mathcal{L}_b[e_2] in
                                let l_2 = copy(live) in
                                   live := l_1; \mathbf{let} \ r_3 = \mathcal{L}_b[e_3] \ \mathbf{in}
                                                    let l_3 = copy(live) in
                                                      live := \max(l_2, l_3); \mathbf{let} \ r = join(r_2, r_3) \ \mathbf{in}
                                                                                    setexs(r, \min(l_2 - l_1, l_3 - l_1)); r
    else if b then \mathcal{L}_b[e_2] else \mathcal{L}_b[e_3]
\mathcal{L}_b\left[\mathbf{let}\ v=e_1\ \mathbf{in}\ e_2\right]=\mathbf{let}\ v=\mathcal{L}_b\left[e_1\right]\ \mathbf{in}
                                    incrc(v); let r = \mathcal{L}_b[e_2] in (gcExcept(v, r); recomputeExs(r); r)
\mathcal{L}_b[f(e_1,\ldots,e_n)] = \mathbf{let} \ r_1 = \mathcal{L}_b[e_1],\ldots,r_n = \mathcal{L}_b[e_n] \ \mathbf{in}
                               incrc(r_1); \ldots; incrc(r_n);
                               let r = f_{Lb}(r_1, \ldots, r_n) in
                                  gcExcept(r_1, r); \dots; gcExcept(r_n, r); recomputeExs(r); r
```

Figure 2: Transformation that produces live heap space-bound functions f_{Lb} . copy copies a vector. + and -, when applied to vectors, denote component-wise sum and difference.

join-values with a primitive branch is uk, irrespective of whether the result of null? on the non-primitive branch is true or false.

When a selector c^{-i} is applied to a join-value j with a primitive branch, it simply aborts by attempting to apply the selector to an arbitrary primitive value. However, if we assume that the given program never applies selectors to primitive values, then the occurrence of $c^{-i}(j)$ in the analysis corresponds to the application of c^{-i} to the non-primitive branch of j in the original program. Some test in a conditional in the original program must prevent c^{-i} from being applied to the primitive branch. Thus, with this assumption, we could use the following alternative definition of c_n^{-i} .

```
\begin{split} c_u^{-i}(v) &= \text{if } v = uk \text{ then } uk \\ & \text{else if } isJoin(v) \\ & \text{then if } length(branches(v)) = 1 \text{ then } c_u^{-i}(first(branches(v))) \\ & \text{else } join(c_u^{-i}(first(branches(v))), c_u^{-i}(second(branches(v)))) \\ & \text{else } c^{-i}(data(v)) \end{split}
```

Applications of selectors to join-values with primitive branches is in fact seen in only one of our examples, namely quicksort. For all of our examples, it is easy to see that for the expected input, selectors are never applied to primitive values. Hence, it is safe to use the above definition of c_u^{-i} for all these examples. In general, a simple static analysis could be used to automatically ascertain such program properties.

4.3 Achieving Tightness of live in the Presence of Join-values

The following example illustrates why live may not be as tight as desired.

let
$$u = cons(1, nil)$$
 in
let $v = cons(2, nil)$ in
(if uk then $cons(3, v)$ else $cons(4, cons(5, u))$)

Let r be the result of the conditional. Let c_i denote the data construction with $cons^{-1}(c_i) = i$. Just after the conditional is evaluated, live includes the sizes of both c_1 and c_2 . live excludes the size of c_3 because the result of the alternative branch containing c_4 and c_5 is larger; so live includes the latter instead of the former. Once v goes out of scope, c_2 is live only through the reference from r. At this point in any execution of the original program, either c_2 and c_3 are live or c_4 and c_5 are live; c_1 is definitely live because of the binding for v. But in the analysis, because of the reference from v, v is kept live and its size is included in live. Thus, join-value v causes live to be loose by one cons.

In general, at any point at which all references to a data construction v are lost except for references from a join-value j, there is a possibility that live is loose because it includes the size of v when it should not. These points arise immediately after decrements to rc(v) caused by (1) a variable or parameter going out of scope or (2) parts of data becoming garbage after the application of a selector. v may then be an excess in live caused by a join-value j which in case (1), is in the result of the function call or the let expression and in case (2), is in the result of the selector. recomputeExs, defined in Figure 3, is called on the results of function calls, let expressions and selectors to compute the exs attributes of join-values in the results and adjust the value of live appropriately.

Observe that v may be a part of a join-value j' that is not in the result of the function call or **let** expression or selector application. It can be shown that loss of references to v at the completion of the function call, **let** expression or selector application, has no effect on exs(j') and so we do not call recomputeExs(j'). This applies to tester applications also. Further, results of testers are boolean and hence may not contain any join-values. So, recomputeExs is not called after tester applications. Note that recomputeExs is used only to obtain tighter bounds, so calling or not calling it at any point in the analysis is safe, i.e., still yields an upper bound on the space usage.

We now formally define the exs attribute of join-values. Consider the stack and live heap as a graph: convalues and join-values in the heap and formal parameters of functions and **let**-bound variables on the stack are vertices; references from variables, con-values and join-values to con-values and join-values, including references in *branches* attributes but excluding references in *joinParents* attributes, are edges. We say that u is *contained-in* v if v is an ancestor of u in every path from a node for a parameter or variable to u. For a

```
join(v_1, v_2) = if \ eq?(v_1, v_2) \ then \ v_1
               else if isPrim(v_1)
                    then if isPrim(v_2) then uk
                           else let result = newjoin([v_2]) in
                                  incrc(v_2); addJoinParent(v_2, result); result
                    else if isPrim(v_2)
                          then let result = newjoin([v_1]) in
                                  incrc(v_1); addJoinParent(v_1, result); result
                          else let result = newjoin([v_1, v_2]) in
                                 incrc(v_1); addJoinParent(v_1, result);
                                 incrc(v_2); addJoinParent(v_2, result);
                                 result
qc(v) = \mathbf{if} \ not(isPrim(v))
         then decrc(v);
               if rc(v) < 0
               then if isJoin(v)
                     then live = live + exs(v);
                            for u in branches(v)
                              delJoinParent(u, v); gc(u)
                     else live[conType(v)] --;
                           for i = 1..arity(v) \ gc(c^{-i}(data(v)))
recomputeExs(v) = \mathbf{if} \ not(isPrim(v)) \ \mathbf{then}
                       if isJoin(v)
                       then if length(branches(v)) = 2 and containedIn_0(branches(v))
                             then let newexs = computeExs(v) in
                                     if newexs > exs(v)
                                     then live = live + exs(v) - newexs; setexs(v, newexs)
                             else for u in branches(v) recomputeExs(u)
                       else for i = 1..arity(v) recomputeExs(c^{-i}(data(v)))
containedIn_0(ls) = if \ null(ls) \ then \ true
                     else if rc(cons^{-1}(ls)) = length(joinParents(cons^{-1}(ls))) = 1
                          then contained In_0(cons^{-2}(ls))
                          else false
```

Figure 3: Auxiliary functions join, gc, recomputeExs and $containedIn_0$. For brevity, we leave out the definition of computeExs which is based on (2) in Section 4.3.

join-value j, let C_j denote the set of all con-values and join-values contained-in j, and let G_j denote the graph comprising vertices and edges reachable from j. A join-path of j is a connected subgraph of G_j containing j and constructed from G_j by selecting at every join-value j' reachable from j, exactly one branch of j' and then, after all selections have been made, eliminating unreachable vertices and edges. Figure 4 contains examples of join-paths. Join-paths of j correspond to data structures represented by j. conCountVec(u) for a con-value u is a constructor count vector in which the count of the constructor type of u is 1 and all other counts are 0. maxJoinPath(j) is the maximum of the sizes of all join-paths of j, where size(P) for a join-path P of j is defined as

$$\mathit{size}(P) = \sum_{u \text{ is a con-value in } P \cap C_j} \mathit{conCountVec}(u)$$

exs(j) is then defined as follows if both branches of j are non-primitive and contained in j (otherwise, exs(j)

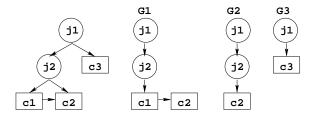


Figure 4: Examples of join-paths. G1, G2 and G3 are join-paths of join-value j1. Circles denote join-values and rectangles denote con-values.

is V_0).

$$exs(j) = total(j) - sub(j) - maxJoinPath(j)$$
 (2)

$$total(j) = \sum_{u \text{ is a convenies in } C} conCountVec(u)$$
 (3)

$$exs(j) = total(j) - sub(j) - maxJoinPath(j)$$

$$total(j) = \sum_{u \text{ is a con-value in } C_j} conCountVec(u)$$

$$sub(j) = \sum_{u \text{ is a join-value in } C_j} exs(u)$$

$$u \text{ is a join-value in } C_j$$

$$(4)$$

The sums and differences of constructor count vectors are computed component-wise. (2) does not result in vectors with negative counts, since sub(j) and maxJoinPath(j) count data in disjoint subsets of C_i . This is justified as follows: if the join-path P which contributes to maxJoinPath(j) contains a join-value j', then P contains a largest join-path of j' and exs(j') counts data in the other join-paths of j'. Hence, exs(j'), for any descendant join-value j' of j, counts data in join-paths that are not part of P. Informally, (2) says "subtract from live everything except a largest join-path and nodes that have already been subtracted from live". In our implementation, exs(j) is computed using (2) if the elements in branches(j) have reference counts equal to 1; exs(j) is conservatively set to V_0 otherwise.

A recomputed value v_2 of exs(j) can be less than the existing value v_1 of exs(j). This happens only if after the computation of v_1 , selectors are applied to j creating a new join-value j' that references parts of j and a subsequent garbage collection leads to the recomputation of exs(j) yielding v_2 . The references from j' to parts of j cause these parts to not be contained in j and so (2) produces a smaller value than the existing exs(i). But selection from i does not alter the fact that only one of the data constructions represented by i is live, so the new smaller value of exs(j) is artificial and is ignored. Figure 5 contains an example of a live heap space-bound function.

Optimizations 5

We use two optimizations that reduce the asymptotic complexity of live heap analysis for many programs. The first optimization avoids calls to recompute Exs on data without join-value descendants. This is done by adding to con-values and join-values a boolean attribute that indicates the presence of join-value descendants.

The second optimization reduces some join-values to con-values, thus avoiding expensive manipulations of the former. At any point p during the execution of a space-bound function, a join-value j with branches b_1 and b_2 and without any join-value descendants may be reduced to b_1 if b_1 leads to equal or greater live heap usage as compared to b_2 . The following discussion also holds with b_1 and b_2 interchanged. Our optimization is conservative and reduces j to b_1 if b_1 and b_2 have the same shape, contain equal primitive values at corresponding locations, and for every descendant d_1 of b_1 that is not contained in j, the corresponding descendant d_2 of b_2 is the same as d_1 . If these conditions are true and additionally, if eq? is not used on j or on data selected from j, then at point p and thereafter, b_1 contributes as much or more to live as compared to b_2 . In the following subsections, we formalize the notion of reducibility of join-values and prove that reduction of join-values is safe, i.e., the optimized space-bound analysis provides upper bounds on live heap usage.

```
reverse(ls, revls) = if null(ls) then revls
                        else reverse(cons^{-2}(ls), cons(cons^{-1}(ls), revls))
reverse_L(ls, revls) = if ( let l = ls in
                                 let l_{null?} = null(data(l)) in
                                  if not(isPrim(l)) and rc(l) = 0 then gc(l);
                          then revls
                          else let arg1 = ( let l = ls in
                                                   let l_{cdr} = cons^{-2}(data(l)) in
                                                     if not(isPrim(l)) and rc(l) = 0
                                                     then gcExcept(l, l_{cdr});
                                                     l_{cdr})^*
                                      arg2 = (live[i_{cons}] + +; maxlive := max(live, maxlive);
                                                 let ls_{car} = \text{ sub-expression} * \text{ with } cons^{-2} \text{ replaced by } cons^{-1}
                                                       revl = revls in
                                                   incrc(ls_{car}); incrc(revl); new(cons(ls_{car}, revl))) in
                                   incrc(arg1); incrc(arg2);
                                   let r = reverse_L(arg1, arg2) in
                                    gcExcept(arg1, r); gcExcept(arg2, r); r
reverse_{L_b}(ls, revls) = let ls_{null?} = (let l = ls in
                                              let l_{null?} = null_u(l) in
                                               if not(isPrim(l)) and rc(l) = 0 then gc(l);
                                               l_{null?}) in
                              if ls_{null?} = uk
                              then let l_1 = copy(live) in
                                       let branch_1 = revls in
                                         let l_2 = copy(live) in
                                           live := l_1;
                                           let branch_2 =
                                                    (let arg1 = ( let l = ls in
                                                                       let l_{cdr} = cons^{-2}u(l) in
                                                                         if not(isPrim(l)) and rc(l) = 0
                                                                         then incrc(l_{cdr}); gc(l); decrc(l_{cdr});
                                                                         l_{cdr})^*
                                                          arg2 = (live[i_{cons}] + +; maxlive := max(live, maxlive);
                                                                     let ls_{car} = \text{sub-expression} * \text{with}
                                                                                     cons^{-2}u replaced by cons^{-1}u
                                                                           revl = revls in
                                                                       incrc(ls_{car}); incrc(revl); new(cons(ls_{car}, revl))) in
                                                     incrc(arg1); incrc(arg2);
                                                     let r = reverse_{L_h}(arg1, arg2) in
                                                       gcExcept(arg1, r); gcExcept(arg2, r); recomputeExs(r); r)^{\dagger} in
                                             let l_3 = copy(live) in
                                               live := \max(l_2, l_3);
                                              let r = join(branch_1, branch_2) in
                                                setexs(r, \min(l_2 - l_1, l_3 - l_1)); r
                              else if ls_{null?}
                                     then revls
                                     else sub-expression †
```

Figure 5: Examples of space and space-bound functions. $reverse_L$ and $reverse_{L_b}$ are the space and space-bound functions, respectively, of reverse.

An edge-ordered rooted directed acyclic graph (DAG) is a tuple $\langle V, E, r \rangle$ where V is a set of vertices, E is a set of edges and r is the root of the DAG, i.e, a node from which all other nodes in the DAG are reached. An edge in E is a tuple $\langle x, y, i \rangle$, where x is the source node, y is the destination node, and i is the index of the edge amongst the out-edges of x.

Recall that the stack and live heap may be viewed as a graph. The subgraph $G_x = \langle V_x, E_x, x \rangle$ comprised of nodes and edges reachable from a node x is an edge-ordered rooted DAG. It is acyclic because we are dealing with a first-order functional language without imperative update. The ordering of fields in data constructions imposes an ordering on the out-edges from nodes. For example, if x is an instance of cons and $cons^{-2}(x) = y$ and y is not a primitive value, then G_x contains the edge $\langle x, y, 2 \rangle$.

Two edge-ordered rooted DAGs $G_1 = \langle V_1, E_1, r_1 \rangle$ and $G_2 = \langle V_2, E_2, r_2 \rangle$ are isomorphic if there exists a one-to-one, onto mapping $f: V_1 \to V_2$ such that $\langle x, y, i \rangle \in E_1$ iff $\langle f(x), f(y), i \rangle \in E_2$, and x and f(x) are data constructions of the same type. If f(x) = y, we say that x corresponds to y and $vice\ versa$.

Reducibility of Join-values. $j = join(b_1, b_2)$ is reducible to b_1 at a point p_0 during execution of a space-bound function if at p_0

R0. j does not have any join-value descendants.

R1. G_{b_1} and G_{b_2} , the DAGs rooted at b_1 and b_2 , are isomorphic.

R2. Corresponding primitive values in b_1 and b_2 and their descendants are equal, taking uk = uk.

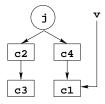
R3. For every node d_1 of G_{b_1} , if d_1 is not contained in j, then d_1 and $f(d_1)$ are the same node.

R0 implies that j represents exactly two data structures: b_1 and b_2 . R1 and R2 state that b_1 and b_2 have the same structure and contents. For example, b_1 and b_2 may be two lists of the same length and containing the same primitive values. The only possible difference between b_1 and b_2 is the particular heap locations they use. No operation in our language can distinguish b_1 and b_2 ; recall that we don't consider eq?. Thus, R1, R2 and R4 ensure that the program's execution is the same regardless of whether b_1 or b_2 is used, except for the heap space used by b_1 and b_2 ; specifically, using b_1 vs. b_2 does not affect any other heap allocations or live heap usage of the program. R3 asserts that b_1 always contributes at least as much to the live heap space as b_2 . For example, this happens if b_2 references data constructions that are live even without references from b_2 and the corresponding data constructions in b_1 are live only because of the references from b_1 .

As an example, consider the following expression.

let
$$v = cons(uk, nil)$$
 in
if uk then $cons(uk, cons(uk, nil))$ else $cons(uk, v)$

The abstract heap at the point just after evaluating the conditional is shown above. Con-values c1 through



c4 are numbered according to the order of creation. The result j of the conditional satisfies conditions R0 through R4, and hence may be reduced to its left branch.

Theorem 1 If $j = join(b_1, b_2)$ is reducible to b_1 at a point p_0 during execution of a space-bound function, then it is safe to replace all references to j with references to b_1 , i.e., in the presence of these replacements the space-bound analysis still returns an upper bound on live heap usage.

Proof: Based on the above arguments, it suffices to show that b_1 contributes at least as much to the live heap space as b_2 .

Let $G_{b_1} = \langle V_{b_1}, E_{b_1}, b_1 \rangle$ and $G_{b_2} = \langle V_{b_2}, E_{b_2}, b_2 \rangle$ be the edge-ordered DAGs rooted at b_1 and b_2 , respectively. Let C_j be the set of nodes contained-in j at p_0 . The contribution of b_i to live is the amount of data contained-in j and referenced by b_i . Observe that data not contained-in j is live regardless of whether a replacement of j with b_1 or b_2 is performed. The contribution of j to live is the maximum of the contributions of b_1 and b_2 . At p_0 ,

contrib. of
$$b_1$$
 to $live = \sum_{x \in (V_{b_1} \bigcap C_j)} conCountVec(x)$ (5)

contrib. of
$$b_2$$
 to $live = \sum_{x \in (V_{b_2} \bigcap C_j)} conCountVec(x)$ (6)

First, we show that (5) is at least as large as (6). R1 states that G_{b_1} and G_{b_2} are isomorphic. Let $f: V_{b_1} \to V_{b_2}$ be the isomorphism. We show that at p_0 , for every node $y \in (V_{b_2} \cap C_j)$, $f^{-1}(y) \in (V_{b_1} \cap C_j)$. By definition, $f^{-1}(y) \in V_{b_1}$. We prove by contradiction that $f^{-1}(y) \in C_j$. Suppose not. Then, according to R3, $f^{-1}(y) = y$. Hence, $y \notin C_j$. This contradicts the assumption that $y \in (V_{b_2} \cap C_j)$, so $f^{-1}(y) \in C_j$. Also, y and $f^{-1}(y)$ are data constructions of the same type. Thus, for each element of the sum in (6), there is a corresponding and equal element of the sum in (5), so (5) is at least as large as (6).

Now, consider a point p_1 after p_0 . If j is dead at p_1 , then the replacement has no more effect on the execution of the space-bound function; the indirect effect through parts of j that were selected out and that are still live, is considered below. Suppose j is live at p_1 . Observe that G_{b_1} and G_{b_2} do not change between p_0 and p_1 , due to the absence of imperative update, and that f is still an isomorphism between G_{b_1} and G_{b_2} . We use C_x^p to refer to the set of nodes contained-in node x at point p; p is omitted when it is clear from context. We show that, for every node $y \in (V_{b_2} \cap C_j^{p_1})$, $f^{-1}(y) \in C_j^{p_1}$. Case 1: $f^{-1}(y) \notin C_j^{p_0}$. Then, according to R3, $f^{-1}(y) = y$. By hypothesis, $y \in C_j^{p_1}$, so $f^{-1}(y) \in C_j^{p_1}$. Case 2: $f^{-1}(y) \in C_j^{p_0}$. We prove by contradiction that $f^{-1}(y) \in C_j^{p_1}$. $f^{-1}(y) \in C_j^{p_0}$ implies that after p_0 , the only way to create references to $f^{-1}(y)$ from outside G_j (the DAG rooted at j) is through selector applications to j. This is thus the only way for $f^{-1}(y)$ to become not contained-in j after p_0 . Since selector applications to j return join-values whose branches are corresponding nodes in V_{b_1} and V_{b_2} , such join-values that reference $f^{-1}(y)$ must also reference y. So, if $f^{-1}(y) \notin C_j$ (as hypothesized for the proof by contradiction), then $y \notin C_j$. This contradicts the assumption $y \in (V_{b_2} \cap C_j^{p_1})$. Hence, $f^{-1}(y) \in C_j^{p_1}$.

We now show that after p_0 , the result of a selector application to b_1 also contributes at least as much to live as compared to the result of the same selector application to b_2 . Let the result of $c_1^{-i_1}(...(c_n^{-i_n}(j)))$ be $j' = join(b'_1, b'_2)$ where $b'_1 \in V_{b_1}$ and $b'_2 \in V_{b_2}$, and let p_1 be a point equal to or after p_0 at which j' is live. At p_1 ,

contrib. of
$$b'_1$$
 to $live = \sum_{x \in (V_{b'_1} \bigcap C_{j'})} conCountVec(x)$ (7)

contrib. of
$$b'_2$$
 to $live = \sum_{x \in (V_{b'_2} \bigcap C_{j'})}^{c_1 \cap C_{j'}} conCountVec(x)$ (8)

Observe that the restriction of f to any subgraph G of G_{b_1} is an isomorphism between G and the corresponding subgraph of G_{b_2} . So, the restriction of f to $G_{b'_1}$ is an isomorphism between $G_{b'_1}$ and $G_{b'_2}$. We refer to this restriction of f as just f. We prove that for every $y \in (V_{b'_2} \cap C_{j'}^{p_1})$, $f^{-1}(y)$ is in $(V_{b'_1} \cap C_{j'}^{p_1})$. We consider two cases. Case 1: $f^{-1}(y) \notin C_j^{p_0}$. Then, $f^{-1}(y) = y$, because of R3. y is in $C_{j'}^{p_1}$, so $f^{-1}(y)$ is in $C_{j'}^{p_1}$. Case 2: $f^{-1}(y) \in C_j^{p_0}$. We prove by contradiction that $f^{-1}(y) \in C_{j'}^{p_1}$. Suppose this is not so. Then, there exist references to $f^{-1}(y)$ from outside G'_j , the DAG rooted at j'. At p_0 , $f^{-1}(y) \in C_j$, so any references to $f^{-1}(y)$ from outside G'_j are created after p_0 , implying p_1 is after (not equal to) p_0 . Furthermore, these references must be due to selector applications to j. Thus, at p_1 , references to $f^{-1}(y)$ from outside $G_{j'}$ are either from j or from the results of selector applications to j. Since both j and results of selector applications to j are

join-values that reference corresponding nodes in V_{b_1} and V_{b_2} , there exist references from outside $G_{j'}$ to y also, so $y \notin C_{j'}^{p_1}$. This contradicts the assumption $y \in (V_{b'_2} \cap C_{j'}^{p_1})$. Hence, $f^{-1}(y)$ is in $C_{j'}^{p_1}$. Thus, for each element of the sum in (8), there is a corresponding and equal element of the sum in (7), so (7) is at least as large as (8).

6 Handling Tail Call Optimization

In some languages, the environment of the caller is discarded only after the completion of all function calls in its body, even if the body contains a function call in tail position. An expression is in *tail position* if it is the last thing that the function does before returning. A *tail call* is a function call in tail position. Tail call optimization [1] allows the caller's environment to be discarded right before executing a tail call. We have extended our analysis to reflect the effect of tail call optimization. In the presence of this optimization, the analysis described in Sections 3 and 4 yields safe but perhaps loose bounds on space usage.

Our extended analysis recognizes function calls in tail position and at the sites of these calls, garbage collects all variables in the current scope. In the original analysis, rc's of arguments are incremented just before function calls and arguments are garbage collected on return from the calls. In the extended analysis, both operations are performed within the function body: increments are done at the start of the function and garbage collection is done just before the result is returned. The latter is performed either just before a tail call or on completion of the last non-call expression in the function body. This structure provides for a simple transformation as well as simpler and more efficient space and bound functions. Figure 6 contains the new transformation that yields space functions. The right hand side uses quasi-quotes '...'; all expressions within quotes, except those of the form " $\mathcal{L}_x[e]$...", are to be taken literally. We make the exception to minimize clutter.

Figures 7, 8 and 9 contain the transformation that yields bound functions. The transformation of conditionals is more involved than \mathcal{L}_b . Consider a conditional (**if** uk **then** e_2 **else** e_3) that is in tail position. Both e_2 and e_3 need to be evaluated. Suppose both e_2 and e_3 contain tail calls. During the evaluation of e_2 , all environment variables u_1, \ldots, u_m are garbage collected just before the tail call. But at the start of evaluation of e_3 , u_1, \ldots, u_m are live, so the effects of the earlier garbage collection have to be reversed before evaluating e_3 . This is done by using recIncre to increment rc's of u_1, \ldots, u_m and all their descendants. Further, during the evaluation of e_3 , u_1, \ldots, u_m are garbage collected just before the tail call in e_3 . Care needs to be taken to ensure that references from the result r_2 of e_2 to u_1, \ldots, u_m do not cause the latter to be counted as live after this garbage collection. So, before evaluating e_3 , the rc's of r_2 and all its descendants are decremented. Similar issues arise when only one of the two branches contain tail calls. Consider the case when neither of the branches contain tail calls. In each branch, u_1, \ldots, u_m are garbage collected after the evaluation of the last non-call expression in the branch. Instead of garbage collecting u_1, \ldots, u_m at the end of e_2 , and again incrementing their rc's and those of their descendants before evaluating e_3 , we simply evaluate e_2 and e_3 as if they were not in tail position and garbage collect u_1, \ldots, u_m just before returning the result of the conditional.

threc? in Figure 9 is used to determine whether or not a given expression contains tail calls. More precisely, given an expression e in tail position, tlrec?(e) determines if all variables in the scope of e are garbage collected in \mathcal{L}_b [e] itself or if they have to be garbage collected after \mathcal{L}_b [e]. This more precise definion is particularly relevant to conditionals. Consider a conditional $e = (\mathbf{if}\ e_1\ \mathbf{then}\ e_2\ \mathbf{else}\ e_3)$ in which one branch, say e_2 , contains a tail call and the other does not. In e_2, u_1, \ldots, u_m are garbage collected before the tail call. l_2 in \mathcal{L}_{xb} [e] is the value of live after the evaluation of e_2 . We need to garbage collect u_1, \ldots, u_m at the end of e_3 in order to ensure that the value of live after the evaluation of e_3 $(l_3\ \text{in}\ \mathcal{L}_{xb}\ [e])$ corresponds to the same "execution point" as l_2 . If neither e_2 nor e_3 contained tail calls it would be sufficient to garbage collect u_1, \ldots, u_m after the entire conditional is evaluated. Therefore, tlrec?(e) is true exactly when either tlrec?(e) or tlrec?(e) is true.

```
f_{L_x}(v_1,\ldots,v_n) = incrc(v_1);\ldots;incrc(v_n);
                          \mathcal{L}_x[e]true
  where e is the body of function f, i.e., f(v_1, \ldots, v_n) = e
\mathcal{L}_x[v] tailpos? = if tailpos? then 'gcExcept(u_1, v); \dots; gcExcept(u_m, v); v' else 'v'
\mathcal{L}_x[l] tailpos? = \mathbf{if} \ tailpos? \mathbf{then} \ `gc(u_1); \dots; gc(u_m); \ l' \ \mathbf{else} \ `l'
\mathcal{L}_x[c(e1,\ldots,en)] tailpos? = 'live[i<sub>c</sub>]++; if (P \cdot live > P \cdot maxlive)
                                                            then for c \in Constructors \ maxlive[i_c] := live[i_c];
                                         let r_1 = {}^{\backprime}\mathcal{L}_x[e_1] false^{\backprime}, {}^{\backprime}\dots, {}^{\backprime}r_n = {}^{\backprime}\mathcal{L}_x[e_n] false^{\backprime} in
                                            incrc(r_1); \ldots; incrc(r_n);
                                           if tailpos?
                                            then 'let r = new(c(r_1, \ldots, r_n)) in
                                                        gcExcept(u_1, r); \ldots; gcExcept(u_m, r); r'
                                           else 'new(c(r_1,\ldots,r_n))'
\mathcal{L}_x[p(e_1,\ldots,e_n)] tailpos? = if tailpos? then 'let r = p(\mathcal{L}_x[e_1] false,\ldots,\mathcal{L}_x[e_n] false) in
                                                                      gc(u_1); \ldots; gc(u_m);
                                         else 'p(\mathcal{L}_x[e_1]false, \dots, \mathcal{L}_x[e_n]false)'
\mathcal{L}_x[c?(e)] tailpos? = 'let x = '\mathcal{L}_x[e] false 'in
                                let r = (if \ isPrim(x) \ then \ c?(x) \ else \ c?(data(x))) \ in
                                   (if not(isPrim(x)) and rc(x) = 0 then gc(x));
                                  if tailpos? then 'gc(u_1); \ldots; gc(u_m); r'
\mathcal{L}_x \left[ c^{-i}(e) \right] tailpos? = \text{`let } x = \text{'} \mathcal{L}_x \left[ e \right] false \text{`in}
                                    let r = c^{-i}(data(x)) in
                                      (if not(isPrim(x)) and rc(x) = 0 then gcExcept(x,r));
                                      if tailpos? then 'gcExcept(u_1, r); \ldots; gcExcept(u_m, r); r'
\mathcal{L}_x [if e_1 then e_2 else e_3] tailpos? = 'if \mathcal{L}_x [e_1] false then \mathcal{L}_x [e_2] tailpos? else \mathcal{L}_x [e_3] tailpos?'
\mathcal{L}_x [let v = e_1 in e_2] tailpos? = 'let v = \mathcal{L}_x [e_1] false in
                                                 if tailpos? then '\mathcal{L}_x[e_2] true'
                                                 else 'let r = \mathcal{L}_x[e_2] false in
                                                            gcExcept(v, r); r'
\mathcal{L}_x[f(e_1,\ldots,e_n)] tailpos? = if tailpos?
                                         then 'let r_1 = \mathcal{L}_x[e_1] false, \dots, r_n = \mathcal{L}_x[e_n] false in
                                                      incrc(r_1); \ldots; incrc(r_n);
                                                      gc(u_1); \ldots; gc(u_m);
                                                      decrc(r_1); \ldots; decrc(r_n);
                                         f_{L_x}(r_1, \dots, r_n)' else 'f_{L_x}(\mathcal{L}_x[e_1] false, \dots, \mathcal{L}_x[e_n] false)'
```

Figure 6: Transformation that produces live heap space functions f_{L_x} handling tail call optimization. u_1 , ..., u_m are variables in the current scope.

7 Experiments

We implemented the analyses and measured the results for several standard list and tree processing programs. Comparisons of results of space functions and bound functions show that bound functions produce accurate

```
f_{L_{rh}}(v_1,\ldots,v_n) = incrc(v_1);\ldots;incrc(v_n);
                                                                     \mathcal{L}_{xb}[e]true'
      where e is the body of function f, i.e., f(v_1, \ldots, v_n) = e
 \mathcal{L}_{xb}[v] tailpos? = \mathbf{if} \ tailpos? \mathbf{then} \ `gcExcept(u_1, v); \dots; gcExcept(u_m, v); \ v` \mathbf{else} \ `v'
 \mathcal{L}_{xb}[l] \ tailpos? = \mathbf{if} \ tailpos? \ \mathbf{then} \ \ `gc(u_1); \dots; gc(u_m); \ l' \ \mathbf{else} \ \ `l'
 \mathcal{L}_{xb}[c(e1,\ldots,en)] tailpos? = `live[i_c] + +; if (P \cdot live > P \cdot maxlive)
                                                                                                                                                   then for c \in Constructors\ maxlive[i_c] := live[i_c];
                                                                                                      let r_1 = {}^{\backprime}\mathcal{L}_{xb}[e_1]false, {}^{\backprime}, {}^{\backprime
                                                                                                             incrc(r_1); \ldots; incrc(r_n);
                                                                                                            if tailpos?
                                                                                                            then 'let r = new(c(r_1, \ldots, r_n)) in
                                                                                                                                            `gcExcept(u_1, r); \dots; gcExcept(u_m, r); recomputeExs(r); r'
                                                                                                            else 'new(c(r_1,\ldots,r_n))'
\mathcal{L}_{xb}[p(e_1,\ldots,e_n)] tailpos? = if tailpos? then 'let r = p_u(\mathcal{L}_{xb}[e_1] false,\ldots,\mathcal{L}_{xb}[e_n] false) in
                                                                                                                                                                             gc(u_1); \ldots; gc(u_m);
r'
                                                                                                      else 'p_u(\mathcal{L}_{xb}[e_1]false, \dots, \mathcal{L}_{xb}[e_n]false)'
p_u(v_1,\ldots,v_n)=\mathbf{if}\ v_1=uk\ or\ \cdots\ or\ v_n=uk\ \mathbf{then}\ uk\ \mathbf{else}\ p(v_1,\ldots,v_n)
 \mathcal{L}_{xb}[c?(e)] tailpos? = 'let x = '\mathcal{L}_{xb}[e] false 'in
                                                                                let r = c?_u(x) in
                                                                                       (if not(isPrim(x)) and rc(x) = 0 then gc(x));
                                                                                      if tailpos? then 'gc(u_1); \ldots; gc(u_m); r'
                                                                                       else 'r'
 c?_u(v) = \mathbf{if} \ v = uk \ \mathbf{then} \ uk
                                  else if isJoin(v) then if length(branches(v)) = 1
                                                                                                                   then join(false, c?_u(first(branches(v))))
                                                                                                                    else join(c?_u(first(branches(v))), c?_u(second(branches(v))))
                                                  else c?(data(v))
\mathcal{L}_{xb} \llbracket c^{-i}(e) \rrbracket \ tailpos? = \text{`let } x = \text{'} \ \mathcal{L}_{xb} \llbracket e \rrbracket \ false \text{`in}
                                                                                         let r = c_u^{-i}(x) in
                                                                                                (if not(isPrim(x)) and rc(x) = 0 then gcExcept(x,r));
                                                                                              if tailpos? then 'gcExcept(u_1, r); \ldots; gcExcept(u_m, r); recomputeExs(r) r'
                                                                                                else 'recomputeExs(r) r'
 c_u^{-i}(v) = \mathbf{if} \ v = uk \ \mathbf{then} \ uk
                                 else if isJoin(v)
                                                  then if length(branches(v)) = 1 then c^{-i}(false)
                                                                        else join(c_u^{-i}(first(branches(v))), c_u^{-i}(second(branches(v))))
                                                  else c^{-i}(data(v))
\mathcal{L}_{xb}\left[\mathbf{let}\ v=e_{1}\ \mathbf{in}\ e_{2}\right]tailpos?='\mathbf{let}\ v=\mathcal{L}_{xb}\left[e_{1}\right]false\ \mathbf{in}
                                                                                                                           incrc(v);
                                                                                                                          if tailpos? then '\mathcal{L}_{xb}[e_2]true'
                                                                                                                          else 'let r = \mathcal{L}_{xb}[e_2] false in
                                                                                                                                                    gcExcept(v,r); recomputeExs(r); r'
\mathcal{L}_{xb}\left[f(e_1,\ldots,e_n)\right] tailpos? = \mathbf{if} \ tailpos?
                                                                                                       then 'let r_1 = \mathcal{L}_{xb}[e_1] false, \dots, r_n = \mathcal{L}_{xb}[e_n] false in
                                                                                                                                    incrc(r_1); \ldots; incrc(r_n);
                                                                                                                                    gc(u_1);\ldots;gc(u_m);
                                                                                                                                    decrc(r_1); \ldots; decrc(r_n);
                                                                                                                                    recomputeExs(r_1); \ldots; recomputeExs(r_n);
                                                                                                                                    f_{L_{xb}}(r_1,\ldots,r_n)
                                                                                                       else 'f_{L_{xb}}(\mathcal{L}_{xb}[e_1]false,\ldots,\mathcal{L}_{xb}[e_n]false)'
```

Figure 7: Transformation that produces live heap space-bound functions $f_{L_{xb}}$ handling tail call optimization.

```
\mathcal{L}_{xb} [if e_1 then e_2 else e_3] tailpos? =
  if tailpos?
  then if tlrec?(e_2)
          then if tlrec?(e_3)
                  then 'let b = \mathcal{L}_{xb}[e_1] false in
                             if b = uk then let l_1 = copy(live) in
                                                     let r_2 = \mathcal{L}_{xb} [e_2] true in
                                                       let l_2 = copy(live) in
                                                         recincrc(u_1); \ldots; recincrc(u_m);
                                                         recdecrc(r_2);
                                                          live := l_1; \mathbf{let} \ r_3 = \mathcal{L}_{xb} [e_3] \ true \ \mathbf{in}
                                                                         let l_3 = copy(live) in
                                                                            live := \max(l_2, l_3);
                                                                            recincrc(r_2); let r = join(r_2, r_3) in
                                                                                                setexs(r, computeExs(r)); r
                             else if b then \mathcal{L}_{xb}[e_2] true else \mathcal{L}_{xb}[e_3] true'
                  else 'let b = \mathcal{L}_{xb}[e_1] false in
                            if b = uk then let l_1 = copy(live) in
                                                    let r_3 = \mathcal{L}_{xb} [e_3] false in
                                                      let l_3 = copy(live) in
                                                        recdecrc(r_3);
                                                        live := l_1; \mathbf{let} \ r_2 = \mathcal{L}_{xb} [e_2] true \mathbf{in}
                                                                        let l_2 = copy(live) in
                                                                          live := \max(l_2, l_3);
                                                                          recincrc(r_3); let r = join(r_2, r_3) in
                                                                                               setexs(r, computeExs(r)); r
                            else if b then \mathcal{L}_{xb}[e_2] true else \mathcal{L}_{xb}[e_3] true
          else if tlrec?(e_3)
                 then 'let b = \mathcal{L}_{xb}[e_1] false in
                            if b = uk then let l_1 = copy(live) in
                                                    let r_2 = \mathcal{L}_{xb} [e_2] false in
                                                      let l_2 = copy(live) in
                                                        recdecrc(r_2);
                                                        live := l_1; \mathbf{let} \ r_3 = \mathcal{L}_{xb} [e_3] true \mathbf{in}
                                                                        let l_2 = copy(live) in
                                                                          live := \max(l_2, l_3);
                                                                          recincrc(r_2); let r = join(r_2, r_3) in
                                                                                               setexs(r, computeExs(r)); r
                            else if b then \mathcal{L}_{xb}[e_2] true else \mathcal{L}_{xb}[e_3] true
                 else 'let b = \mathcal{L}_{xb}[e_1] false in
                           if b = uk then let l_1 = copy(live) in
                                                   let r_2 = \mathcal{L}_{xb}[e_2] false in
                                                     let l_2 = copy(live) in
                                                       live := l_1; \mathbf{let} \ r_3 = \mathcal{L}_{xb} [e_3] false \mathbf{in}
                                                                       let l_3 = copy(live) in
                                                                         live := \max(l_2, l_3);
                                                                         let r = join(r_2, r_3) in
                                                                           setexs(r, \min(l_2 - l_1, l_3 - l_1));
                                                                           gcExcept(u_1, r); \ldots; gcExcept(u_m, r);
                          else if b then \mathcal{L}_{xb}[e_2]true else \mathcal{L}_{xb}[e_3]true
  else 'let b = \mathcal{L}_{xb}[e_1] false in
           if b = uk then let l_1 = copy(live) in
                                   let r_2 = \mathcal{L}_{xb}[e_2] false in
                                     let l_2 = copy(live) in
                                        live := l_1; \mathbf{let} \ r_3 = \mathcal{L}_{xb} [e_3] false \mathbf{in}
                                                        let l_3 = copy(live) in
                                                          live := \max(l_2, l_3); \mathbf{let} \ r = join(r_2, r_3) \ \mathbf{in}
                                                                                      setexs(r, \min(l_2 - l_1, l_3 - l_1)); r
```

```
recincrc(v) = incrc(v);
                if (rc(v) \leq 1)
                then if isJoin(v)
                       then for u in branches(v) recincrec(u)
                       else for i = 1..arity(v) \ recincrc(c^{-i}(data(v)))
recdecrc(v) = decrc(v);
                if (rc(v) \leq 0)
                then if isJoin(v)
                       then for u in branches(v) recdecrc(u)
                       else for i = 1..arity(v) \ recdecrc(c^{-i}(data(v)))
tlrec?(e) = \mathbf{case}\ e\ \mathbf{of}
                                              false
                                              false
                                           : false
                p(e_1, ..., e_n)
                c(e_1, ..., e_n)
                                          : false
                c?(e)
                                              false
                c^{-i}(e)
                                           : false
                if e_1 then e_2 else e_3
                                         : tlrec?(e_2) \lor tlrec?(e_3)
                let v = e_1 in e_2
                                           : tlrec?(e_2)
                f(e_1, ..., e_n)
                                              true
```

Figure 9: Auxiliary functions recincre, recdecre and three?.

results (i.e., tight bounds) for all these examples. The results are consistent with the expected asymptotic space complexities of the programs. We measured the running times of space and bound functions of all examples. For most of the examples, the bound functions have the same asymptotic time complexities as the corresponding space functions. For all examples, a comparison of the running times of bound functions and the running times of space functions multiplied by the number of represented inputs showed that the bound functions are asymptotically faster than applying the corresponding space functions to all represented inputs. The non-termination issue mentioned in Section 1 is not a problem for any of these examples.

Figure 10 contains the results of live heap space analysis on some examples. For all examples except quicksort, we show only the results of bound functions on partially known inputs, because they are the same as the results of the space functions on worst-case input. Reversal using append is the standard quadratic-time version. The version of merge sort tested is the one that splits the input list into sublists containing the elements at odd and even positions. Dynamic programming algorithms [5] are used for binomial coefficient, longest common subsequence and string edit. Binary-tree insertion involves insertion of an item into a complete binary tree in which each node is a list containing an element and left and right subtrees.

The partially known inputs for the bound functions of reversal and sorting are lists of known lengths n where all elements are uk; those for longest common subsequence and string edit are two such lists of equal length n. The bound function for binary-tree insertion inserts uk into a complete binary tree of known height h with unknown elements. For binomial coefficient we use integer arguments, n and n-2, since it was found that for a given n, a value of n-2 for the second argument leads to maximum live heap usage.

The difference between the results of the space and bound functions of quicksort is explained as follows. Quicksort selects a pivot element p from the given list ls and splits the list into two lists: one containing all elements of ls that are lesser than or equal to p and the other containing elements that are greater than p. Suppose ls has size n. Since all elements in ls are uk, the bound function incorrectly concludes that each of the two lists has worst-case size n-1. In reality, the sum of the sizes of the two lists is n-1. Further, since quicksort is called recursively on the two lists, inaccuracies at every recursion level quickly add up resulting in the exponential growth of the bound function results.

In time analysis [22] and stack space analysis [31], the bound functions of quicksort run into the non-

termination problem. This is because results of conditionals are summarized into partially known structures and the recursion in quicksort depends on an unknown aspect of such a partially known structure. In live heap analysis, we retain more information about the results of branches of conditionals by using join-values rather than partially known structures. The mentioned recursion in quicksort now has sufficient information to terminate.

,	ersal opend		rtion ort		ection sort		nerge sort				t coefficient		subseq.		string edit		1	binary tr insert	
n	result	n	result	n	result	\overline{n}	result	\overline{n}	space	bound	n	result	n	result	n	result	h	n	result
50	149	50	149	50	1325	2	5	1	1	1	50	194	50	202	50	500	1	3	18
100	299	100	299	100	5150	5	16	2	6	6	100	394	100	402	100	1000	2	7	33
250	749	250	749	250	31625	10	31	5	24	51	250	994	250	1002	250	2500	4	31	111
500	1499	500	1499	500	125750	12	36	6	32	100	500	1994	500	2002	500	5000	7	255	792
1000	2999	1000	2999	1000	501500	15	45	7	41	197	1000	3994	1000	4002	1000	10000	9	1023	3102

Figure 10: Results of live heap space analysis. n is the input size except in the case of binomial coefficient, in which n is the first argument. For binary tree insert, h is the height of the complete binary tree and n is the number of nodes in the tree.

	reversal	w/app	end		insert	ion sor	t	selection sort					merge sort			
n	S	В	Bopt	n	S	В	Bopt	n	S	В	Bopt	n	S	В	Bopt	
10^{1}	1.0 m	0	1.0 m	10^{1}	1.0 m	1.0 M	9.0 m	10^{1}	0	37.3 s	10.0 m	10^{1}	2.0 m	$0.9 \mathrm{\ s}$	$0.7 \mathrm{\ s}$	
10^{2}	$0.1 \mathrm{\ s}$	$0.3 \mathrm{\ s}$	$0.1 \mathrm{\ s}$	10^{2}	$0.1 \mathrm{\ s}$		$5.1 \mathrm{\ s}$	10^{2}	$0.2 \mathrm{\ s}$				40.0 m			
10^{3}	$10.7 \mathrm{\ s}$	$3.5~\mathrm{M}$	$11.9 \mathrm{s}$	10^{3}	$12.8 \mathrm{\ s}$		$1.5~\mathrm{H}$	10^{3}	27.3 s		1.5 H	10^{3}	$0.7 \mathrm{\ s}$			

ŀ	oinomial	coeffic	cient	long	gest con	nmon	subseq.	string edit					binary tree insert					
n	S	В	Bopt	n	S	В	Bopt	n	S	В	Bopt	h	n	S	В	Bopt		
10^{1}	5.0 m	1.0 m	5.0 m	10^{1}	10.0 m	$0.8 \mathrm{\ s}$	44.0 m	10^{1}	15.0 m	25.0 m	$20.0 \mathrm{m}$	2	7	0	3.0 m	3.0 m		
10^{2}	$0.1 \mathrm{\ s}$	$0.4 \mathrm{\ s}$	$0.1 \mathrm{\ s}$	10^{2}	$6.6 \mathrm{\ s}$		$24.0 \ s$	10^{2}	$7.1 \mathrm{\ s}$	$13.5 \mathrm{\ s}$	$7.4 \mathrm{\ s}$	6	127	4.0 m	$0.1 \mathrm{\ s}$	$0.1 \mathrm{\ s}$		
10^{3}	$11.6 \mathrm{\ s}$	$4.7~\mathrm{M}$	$11.6 \mathrm{\ s}$	10^{3}	2.0 H		7.1 H	10^{3}	2.2 H	3.7 H	$2.3~\mathrm{H}$	9	1023	34.0 m	$1.2 \mathrm{\ s}$	$1.2 \mathrm{\ s}$		

Figure 11: Running times of live heap space and live heap space-bound functions. Columns S, B and Bopt contain times of space, unoptimized space-bound and optimized space-bound functions, respectively. n and h are as in Figure 10. m is milliseconds, s is seconds, M is minutes and H is hours. Blank fields in B columns indicate analyses that were aborted after 7 days. Blank fields in Bopt columns indicate analyses that were aborted after 2 days.

The results in Figure 10 include the space used by top-level arguments since these arguments are indeed live throughout the execution of the program. Figure 11 contains running times of live heap space analysis on a sampling of input sizes. For all examples, the live heap space function has the same asymptotic time complexity as the original function. The time complexities of the live heap space-bound functions of reverse using append, binomial coefficient, string edit and longest common subsequence are the same as the complexities of the corresponding original functions. The time complexities of the bound functions of insertion sort and selection sort are a linear factor more than those of the original functions. The linear factor is due to the computation involved in the reduction of join-values. The running time of the bound function of merge sort is more than polynomial in the size of the input. This is because the analysis examines all $(n+m)!/(n! \times m!)$ ways in which two sorted lists of sizes n and m may be merged in sorted order. The running time of the bound function of binary tree insert is polynomial in the size of the input.

The first optimization in Section 5 improves the asymptotic complexities of reverse using append and binomial coefficient by a linear factor. The second optimization improves the asymptotic complexities of insertion sort, selection sort and longest common subsequence from greater than polynomial to polynomial. These speedups are shown in Figure 11.

Figure 12 shows the results of the space analysis that deals with tail call optimization. The corresponding space-bound analysis is not yet implemented. List reversal is the standard linear-time version. The optimized Ackermann function example [23] is a systematically derived program which has much better time complexity

than the classical version. In all of these examples, the main function is tail recursive. The inputs for the first three examples are known lists of length n that lead to the worst-case space usage. The inputs for Ackermann's function are known integers, n and m. The space complexity of this program was worked out by hand to be O(n), but it is hard to see this because of the complicated space usage of the program. For each value of n for Ackermann's function in Figure 12, all values of the second argument in the range [2, 10] produced the same space usage. While this does not prove O(n) space usage, it does help confirm that, for a given n, the space usage is independent of m. Computing Ackermann's function for n > 3 is famously expensive.

tail r	ecursive	tail	recursive	tail 1	recursive	optimized				
list 1	reversal	inser	tion sort	selec	tion sort	Α	Ackermann			
n	result	n	result	n	result	n	result			
50	51	50	100	50	100	0	2			
100	101	100	200	100	200	1	8			
250	251	250	500	250	500	2	15			
500	501	500	1000	500	1000	3	25			
1000	1001	1000	2000	1000	2000					

Figure 12: Results of live heap space functions handling tail call optimization. n is the input size except in the case of optimized Ackermann function, in which n is the first argument; the second argument m ranges over the interval [2, 10], for each n.

Comparing the results in Figure 12 with the space usage of the corresponding non-tail-recursive programs, shown in Figure 10, we see that the tail recursive versions use less live heap space. For example, tail-recursive selection sort uses $O(n^2)$ space. We also applied our extended analysis to the examples of Figure 10. The results were either the same or differed by a constant amount of 1 or 2 cons cells. The differences occur because the programs do contain some function calls in tail position.

Figure 13 contains closed forms or recurrence relations for the live heap space used by the examples. We obtained closed forms by using Matlab to fit polynomials, wherever applicable, to the data in Figure 10. The recurrence relation for merge sort and the closed form for binary tree insert were obtained by hand from the results in Figure 10.

tail recursive reversal	n+1
reversal w/append	0, n = 0 3n - 1, n > 0
	0, n = 0
insertion sort	3n - 1, n > 0
tail rec. insertion sort	2n, n > 0
selection sort	$\frac{n(n+3)}{2}$
tail rec. selection sort	2n
merge sort	$n, n = 0, 1$ $S(\lceil \frac{n}{2} \rceil) + 2n - \lceil \frac{n}{2} \rceil, n > 1$

binomial coefficient	1, m = 0, n 2m + 2, m = n - 1 4m + 2, otherwise
longest common subseq.	n + m + 1, n = 0 or $m = 02m + 3, n = 1$ and $m > 0n + 3m + 2$, otherwise
string edit	3n + 3m + 1, n = 0 or $m = 05n + 5m$, otherwise
binary tree insert	$3(2^{h+1} + h + 1)$

Figure 13: Closed forms or recurrence relations for live heap space used by example programs. n and m are the sizes of the first and second (if any) arguments. For binomial coefficient, n and m are themselves the first and second arguments. For binary tree insert, h is the height of the complete binary tree.

We applied our analysis to a 600-line calendar benchmark. The partially known inputs used are partially known dates. The analysis takes only a few seconds to complete and yields tight bounds, providing preliminary evidence for the scalability of our method. We plan to analyze more benchmarks. We have also used the analysis in teaching programming languages courses.

8 Discussion

Scalability. For large programs or programs with sophisticated control structures, the analysis is efficient if the input parameters are small, but for larger parameters, efficiency might be a challenge. However, from the results of bound analysis on smaller inputs, we may semi-automatically derive closed forms and/or recurrence relations that describe the program's space usage, by fitting a given functional form to the analysis results. Also, when closed forms or recurrence relations are known, we may use the results of the analysis to determine exact coefficients. The closed forms or recurrence relations may then be used to determine space bounds for large inputs.

Termination. The space function terminates iff the original program terminates. The bound function might not terminate, even when the original program does if the recursive structure of the original program directly or indirectly depends on unknown parts of a partially known input structure. For example, if the given partially known input structure is uk, then the bound function for any recursive program does not terminate; if such a bound function counts new space, then the original program might indeed take an unbounded amount of space. Indirect dependency on unknown data can be caused by an imprecise join operation. Making the join operation more precise eliminates this source of non-termination.

Although there are methods to deal with non-termination, incorporating such methods in our analysis could result in loose bounds on space usage, even for programs for which non-termination is not a problem. Further, non-termination is not a problem in any of the examples we analyzed.

Inputs to Bound Functions. To analyze space usage with respect to some property of the input, we need to formulate sets of partially known inputs that represent all actual inputs with that characteristic, e.g., all lists with length n, all binary trees of height h or all binary trees with n nodes. As an example, $\{(uk,(uk,nil,nil),(uk,nil,(uk,nil,nil)),(uk,nil,nil),(uk,nil,nil),(uk,nil,nil)\}$ represents all binary trees of height 1, each node being a list of the element and left and right subtrees. Often, formulating such sets of partially known inputs is straightforward but tedious for the user to do by hand. However, it is easy to write programs that generate such sets of partially known inputs.

Imperative Update and Higher-Order Functions. The ideas in this paper may be combined with reference-counting garbage collection extended to handle cycles [3] or with other garbage collection algorithms, such as mark and sweep, to obtain a live heap space analysis for imperative languages. They may also be combined with techniques for analysis of higher-order functions [30, 11].

Correctness A complete proof of correctness of the analysis is yet to be formulated. Included below is one part of the proof.

Lemma 1 For a join-value j, it is safe to ignore a smaller recomputed value of exs(j).

Proof: Suppose exs(j) is computed at point p and then recomputed at point q and exs(j) at q is less than exs(j) at p. We prove that this may happen only if after p, a selector is applied to j, thus creating a new join-value j' which references parts of j. The referenced parts of j are no longer contained-in j and when exs(j) is recomputed at q, it is smaller than the earlier value of exs(j). It is indeed safe to ignore any decrease in exs(j) caused by selector application because selection does not alter the fact that only one of the data constructions represented by j is live. We keep only one such data construction live by recording the excess in the exs field of j and setting exs(j'), for any j' selected from j, to V_0 . This ensures that excesses are treated as such exactly once. From the definition of exs(2), exs(j) decreases only if one or more of the following occurs:

1. total(j) decreases: this happens if some convalue $c \in C_j^p$ is not contained in j at q. Since c is contained in j at p, there is no way to create a new reference to c other than by applying a selector to j. As we explained earlier, we may safely ignore decreases of exs(j) caused by selection from j.

- 2. max(j) increases: this can happen only if some con-value $c \notin C_j^p$ is contained-in j at q and c is in a maximum-sized join-path of j at q. By definition, the size of c is also included in total(j) at q. Thus, c being contained-in j at q (and not p) causes at least as much increase in total(j) as in max(j). Thus, an increase in max(j) doesn't actually contribute to a decrease in exs(j).
- 3. sub(j) increases : sub(j) increases only if (a) for some join-value $j' \in C_j^p \cap C_j^q$, exs(j') increases (after p) or (b) there exists a join-value $j' \in C_j^q / C_j^p$ such that $exs(j') > V_0$.
 - (a) $j' \in C_j^p$ implies that from point p onwards, it is impossible to obtain a direct reference to j', i.e., no expression may evaluate to j'. Expressions may evaluate to results that contain j' but even then every path from such a result r to j' must contain a join-value, either j itself or a join-value obtained by selection from j. The analysis calls recomputeExs only on results of expressions. Thus, after point p, recomputeExs is never called directly on j'. It may be called on a result r that satisfies the constraints just described. By definition, recomputeExs(r) recomputes the exs fields of any join-value ancestors of j' but does not recompute exs(j'). Hence, for all $j' \in C_j^p$, exs(j') at p' = exs(j') at p, for any point p' > p. (3a) does not cause any increase in sub(j).
 - (b) Let j' be a join-value in $C_j^q \setminus C_j^p$ such that exs(j') at $q > V_0$. exs(j') includes only the sizes of convalues contained-in j'. Since j' is contained-in j, any con-value $c \in C_j'^q$ is also contained-in j at q. Therefore, total(j) at q also includes the size of c. Thus, any increase in sub(j) is accompanied by at least an equal increase in total(j), effectively leaving exs(j) unchanged or larger than before.

9 Related Work

There has been a large amount of work on analyzing program cost or resource complexities, but the majority of it is on time analysis, e.g., [21, 28, 30, 22]. Stack space and heap allocation analysis [31] is similar to time analysis [22]. Analysis of live heap space is different because it involves explicit analysis of the graph structure of the data.

Most of the work related to analysis of space is on analysis of cache behavior, e.g., [32, 10], much of which is at a lower language level, for compiler generated code, while our analyses are at source level and can serve many purposes, as discussed in Section 1. Live heap space analysis is also a first step towards analyzing cache behavior in the presence of garbage collection.

Persson's work on live memory analysis [26] for an object-oriented language requires programmers to give annotations, including specific numbers as bounds for the size of recursive data structures. His work is preliminary: the presentation is informal, with a few formulas summarizing sizes of data in bytes based on the annotations, and only one example, summing a list, is given. Our analysis is able to compute bounds based on input size only, without program annotations.

Unlike static reference counting used in analysis for compile-time garbage collection [18, 16], our analysis uses a reference counting method similar to that in run-time garbage collection. While the former keeps track of pointers to memory cells that will be used later in the execution, the latter maintains pointers reachable from the stack at the current point in execution. Our analysis could be modified so that decrc(v) is called when a parameter or **let**-variable won't be used again (instead of waiting until v goes out of scope). Our current analysis corresponds to the garbage collection behavior in, e.g., JVMs from Sun, IBM, and Transvirtual. Inoue and others [15] analyze functional programs to detect run-time garbage conservatively at compile-time. Their result is an approximation without any information about the input. Also, they do not compute the size of live space.

Several type systems [14, 13, 6] have been proposed for reasoning about space and time bounds, and some of them include implementations of type checkers [14, 6]. They require programmers to annotate their programs with cost functions as types. Furthermore, some programs must be rewritten to have feasible types [14, 13].

Chin and Khoo [4] propose a method for calculating sized types by inferring constraints on size and then simplifying the constraints using Omega [27]. Their analysis results do not correspond to live heap space in general. Further, Omega can only reason about constraints expressed as linear functions.

To summarize, this work is a first attempt to analyze live heap space automatically and accurately using source-level program analysis and transformations. The analysis can be modified to reflect the effect of optimization of tail recursion. The ideas in this paper may be combined with reference-counting garbage collection extended to handle cycles [3] or with other garbage collection algorithms, such as mark and sweep, to obtain a live heap space analysis for imperative languages. They may also be combined with techniques for analysis of higher-order functions [30, 22].

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