R's .Call interface

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1 Example: composite_linkage_disequilibrium

As a simple illustration of the .Call interface, the following function produces a wrapper around the composite linkage disequilibrium calculation introduced in the discussion of .C. The function takes R objects (rather than just the C representation of the data in the object; lines 1, 2), checks that the objects are of the right type (lines 5-12), extracts relevant information (e.g., the dimensions of the matrix of SNPs; lines 15-17), allocates memory for the result (lines 20-21), calls the original C function (lines 24-27), and arranges to clean up and return the result (lines 29, 30).

```
SEXP
1
   composite_linkage_disequilibrium_1 (SEXP snp_r, SEXP width_r)
2
3
   {
        /* check incoming types */
4
        if (!IS_INTEGER(width_r) ||
5
            LENGTH(width_r) != 1 ||
6
            INTEGER(width_r)[0] == NA_INTEGER)
7
8
        {
            error ("'width' must be a single non-NA integer");
9
10
        }
        if (!IS_RAW(snp_r) || LENGTH(GET_DIM(snp_r)) != 2)
11
            error("'snp' must be matrix of raw()");
12
13
        /* retrieve inputs */
14
        int width = INTEGER(width_r)[0],
15
            n_{sub} = INTEGER(GET_DIM(snp_r))[0],
16
17
            n_{snp} = INTEGER(GET_DIM(snp_r))[1];
18
        /* allocate memory for return */
19
       SEXP delta_r:
20
       PROTECT(delta_r = allocMatrix(REALSXP, width, n_snp - width));
21
22
        /* do the calculation */
23
        composite_linkage_disequilibrium (RAW( snp_r ),
24
25
                                           &n_sub, &n_snp, &width,
```

```
REAL(delta_r));
```

26

2 Key components

S-Expressions R objects like numeric() or integer() are actually represented at the C level as a data structure called an *S-expression*, or SEXP. These are defined in the file R_HOME/include/Rinternals.h. They consist of book-keeping information (e.g., what type of data the SEXP holds, and whether there are 0, 1 or more symbols in the R session that have referred to the SEXP) as well as the actual data (e..g, vector of double or int values) associated with the SEXP.

lines 1-2 of the source code define a typical .Call entry point: the function takes two SEXP representing R objects, and returns a SEXP representing an R object.

Interface to R Internals The SEXP are more like R objects than the pointers we've seen in the .C entry point, and in particular they can be queried for properties such as their type (e.g., IS_INTEGER) or length (LENGTH, as in lines 5 and 6. R actually defines several different sets of functions for accessing objects at the C level. These interfaces are defined in R_HOME/include/Rdefines.h and R_HOME/include/Rdefines.h. Looking in these files, you'll see that the length of an object can be determined in a number of ways, including LENGTH, GET_LENGTH, length, and Rf_length. For technical reasons involving name resolution it is probably best to use the Rf_* interface, but in practice packages adopt different approaches.

It is informative to follow functions in either interface to their implementation, typically by checking out the R source code. This can be a lot of fun! For instance, Rf_length is defined as a function at Rinternals.h:1047, and implemented as an inline function at Rinlinedfuns.h:86, where for many types of SEXP it calls the *macro* LENGTH defined at Rinternals.h:267. Conversely, the *function* LENGTH at Rinternals.h:379 is what R package code sees; it too is redefined to point to the inline implementation at Rinlinedfuns.h:86. And finally, in an R session the definition of length is

> length
function (x) .Primitive("length")

R maps this to a C level function called do_length in a table in src/main/names.c:196. This function is defined in src/main/array.c:393, and eventually arrives at our old friend the inlined function at Rinlinedfuns.h:86.

Accessing data contained in R objects Several functions provide access to the data contained in R objects. We see this in lines 7 and 15-17, where the INTEGER function is used to retrieve a C int* pointer. Note that R uses 1-based indexing, but in C the same element is now at position 0. The C level interface provides some convenience functions, such as GET_DIM for retrieving the dimensions of an object that is a matrix. This actually returns a SEXP, which can be queried (as on line 11) for its length.

Allocating memory In R, one can create an object (e.g., numeric(10)) and not worry about what happens to the memory that is used to represent that object. One can ask R to allocate memory using a function like allocMatrix on line 21. This returns an SEXP containing enough room for the requested data, in this case an array of doubles.

R is managing memory, and in particular the allocMatrix function has asked R to move a portion of memory from its pool of available memory to its pool of memory that is in use. R periodically runs a garbage collector to see if any memory that is currently in use can be reclaimed for the pool of memory that can be allocated again. R does this by determining whether symbols in an R session exist that point to each block of memory. For instance, x <- integer(10) associates a portion of R's memory with the variable x. When the garbage collector runs, it notes that there is a symbol x and that it references memory that cannot be garbage collected. If the user says rm(x), then the next time the garbage collector runs it will note that the memory once pointed to by x is not referenced by any current symbol, and will mark the memory as suitable for garbage collection.

Line 21 allocates memory and assigns it to a C variable. But the R garbage collector does not know about C variables, so if the garbage collector were to run, it would think the memory could be collected. This would be very bad – the memory we thought we had access to could instead be assigned and manipulated by some other C level variable. We tell R not to garbage collect memory that has been allocated at the C level but has not been assigned to an R symbol with the **PROTECT** command, as on line 21. This puts the allocated **SEXP** on a protection stack that the garbage collector consults before moving memory from its in-use to its free memory pool.

Notice in line 29 that we call UNPROTECT. This removes the most recently added SEXP from the protection stack, making it available for garbage collection. Immediately after this, and before the garbage collector has a chance to run, we return the allocated memory from our .Call function to the R level. If it is assigned to an R-level variable, then it is again safe from garbage collection. If it is not assigned to a variable, it will be garbage collected (so we will not leak memory).

R ensures that the protection stack is in the same state when it returns from a .Call as when it entered it, so functions like allocMatrix do not result in 'memory leaks'; the more common problem is that the user forgets to PROTECT some allocated memory, the code functions properly most of the time, but once in a while the garbage collector runs mid-way through a .Call and surprising things happen. These issues can be difficult to track down.

R has additional ways in which memory can be requested. This is typically used when one wants to represent native C data types rather than R objects. This memory is allocated with R_alloc or Calloc, defined in $R_HOME/include/R_ext/Memory.h$ and RS.h. Memory allocated with R_alloc is automatically released when R returns from .Call. In contrast, memory allocated with Calloc is not explicitly managed by R, we are responsible for releasing the memory when we are done with it, using the Free function. This is useful when we want to allocate memory at the C level, and use it across several calls from R. For instance, the function that opened a SQLite data base connection allocated memory to represent the connection, and managed this across several transitions between the R and C levels.

3 Directions and Resources

The "Writing R Extensions" manual has several sections on the .Call interface, including calling back in to R either for mathematical functions or to evaluate arbitrary R code. Chamber's Software for Data Analysis and Gentleman's R Programming for Bioinformatics include sections on calls to native languages.