

# Lecture: Interfaces to C (and other languages)

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# Interfaces to C (and other languages)

## When to interface with C?

- ▶ Access facilities available in other libraries, e.g., Rgraphviz and the `graphviz` library for graph manipulation and visualization.
- ▶ Perform algorithms not easily captured by the R programming model, e.g., efficiently updating 'sliding window' statistics (??)
- ▶ Represent data structures that do not fit easily into the R programming model, e.g., Biostrings maintains a single 'read-only' copy of a (e.g., nucleotide) sequence, with light-weight 'views'.
- ▶ Move slow R computations to C, for speed.

# Tools for assessing code I

`codetools` Identify likely programming errors, e.g., referencing global variables.

`system.time` Overall time evaluating code chunk.

`Rprof` Profile time spent in each R function (and optionally memory-allocations).

`tracemem` Record when an object is 'duplicated'.

`gctorture` Flags internal (C-level) errors related to R memory management.

`valgrind` C-level user memory management errors.

# Tools for assessing code II

## Caveats:

- ▶ Memory information (from Rprofmem) requires R to be built with memory profiling enabled
  - ▶ Linux: `$R_SRC/configure --enable-memory-profiling`
  - ▶ Windows:  
`cd $R_SRC/src/gnuwin32`  
`make R_MEMORY_PROFILING=T`
- ▶ `valgrind` is a linux-only tool.

## Useful to remember:

- ▶ *Copy on change* semantics describe (approximately) how R manages memory.

## Understand R's use of memory: copy-on-change I

```
> v <- list(5)
> tracemem(v)

[1] "<0x01cc1c80>"
```

```
> v[[1]] <- 1
> w <- v
> v[[1]] <- 1
```

```
tracemem[0x01cc1c80 -> 0x01a98350]: eval.with.vis doTryCatch
```

```
> v <- list(5)
> tracemem(v)

[1] "<0x01358e88>"
```

## Understand R's use of memory: copy-on-change II

```
> f <- function(x) {  
+   x[[1]] <- 1  
+   x  
+ }  
> v <- f(v)
```

```
tracemem[0x01358e88 -> 0x01d7dd08]: f eval.with.vis doTryCa
```

```
> v <- list(5)  
> tracemem(v)
```

```
[1] "<0x01d81bc8>"
```

```
> g <- function(x) x[[1]]  
> g(v)
```

```
[1] 5
```

# Think in R

E.g., for loops imply memory copies.

```
> reps <- 10000
> system.time({
+   r1 <- list()
+   for (i in seq(1, reps)) r1[[i]] <- i
+ }, gcFirst = TRUE)[["user.self"]]

[1] 3.11
```

```
> system.time({
+   r2 <- lapply(seq(1, reps), function(i) i)
+ }, gcFirst = TRUE)[["user.self"]]

[1] 0.05
```

## Use existing methods

E.g., running median

```
> v <- seq(1, reps)
> window <- 10
> system.time({
+   r1 <- sapply(seq(1, length(v) - window),
+               function(i, v, w) median(v[seq(i,
+               i + w)]), v = v, w = window)
+ }, gcFirst = TRUE)[["user.self"]]
```

```
[1] 4
```

```
> system.time({
+   r2 <- runmed(v, window + 1)
+ }, gcFirst = TRUE)[["user.self"]]
```

```
[1] 0
```



## Understand Rprof I

```
> f <- function(reps) g(reps)
> g <- function(reps) {
+   lst <- list()
+   for (i in seq(1, reps)) lst[[i]] = i
+   lst
+ }
> Rprof()
> res <- f(10000)
> Rprof(NULL)
```

## Understand Rprof II

```
> summaryRprof()$by.self[c("f", "g"), ]  
  
  self.time self.pct total.time total.pct  
f      0.00      0      2.06      99  
g      2.06     99      2.06      99
```

- ▶ `self.time`: time spent *in* the function.
- ▶ `total.time`: time spent in the function, and all contained functions.
- ▶ Most effective when profiling small pieces of code.

# Interfaces to C

## .C

- ▶ R types coerced to familiar C types
- ▶ Useful for quick algorithms or simple interfaces to existing libraries.

## .Call

- ▶ C-level access to R data structures and language features.
- ▶ Much greater knowledge and responsibility.
- ▶ Useful for calling R functions from C code, manipulating R objects, developing extensive library interfaces, implementing novel data structures.

## Also

- ▶ .Fortran: calling Fortran code
- ▶ .Internal, .External: primarily useful for understanding R.
- ▶ Main references: *Writing R Extensions*, *R internals*.

# Overall approach

## R-level

- ▶ Write a wrapper function to perform error checking and invoke C code.
- ▶ Load shared library into R using `dyn.load`.
- ▶ Invoke from R using `.C` or `.Call`.

## C- and system-level

- ▶ Write C functions, with argument and return types depending on whether function will use `.C` or `.Call`.
- ▶ Compile as a shared library, using shell script `R CMD SHLIB`.
- ▶ `Makefile` and `Makevars` often *not* necessary.

## Incorporating C code into packages.

- ▶ Add code to package `src` directory.
- ▶ Write C code to 'register' native routines.
- ▶ Load with `useDynLib` in `NAMESPACE` file.

## The .C interface

```
> noquote(names(as.list(args(.C))))
```

```
[1] name      ...      NAOK      DUP      PACKAGE  
[6] ENCODING
```

*name* Name of C routine.

*...* Arguments to *name*, in order (names are *not* matched to C argument names).

*NAOK* Allow NA and other special values to be passed to C.

*DUP* Duplicate the object to be passed to C? Almost always a good idea to do this.

*PACKAGE* Look for *name* in the dll of the specified package.

*ENCODING* Encoding used for character vectors.

Return value is a (possibly named) list corresponding to ....

## .C example I

For each element  $x[i]$ , find the minimum of  $x[i]-y[j]$  over all  $j$ . E.g., in file R/distance.R

```
> minimumDistance <- function(x, y) {  
+   xlen <- length(x)  
+   ylen <- length(y)  
+   if (ylen < 2)  
+     return(x - rep(y, xlen))  
+   .C("min_dist", as.numeric(x), xlen, as.numeric(y),  
+     ylen, dist = numeric(xlen))$dist  
+ }
```

## .C example II

E.g., in file src/dist.c

```
#include <R.h>

void min_dist(double *x, int *x_len,
              double *y, int *y_len, double *dist) {
    int i, j;
    double cur;
    for (i = 0; i < *x_len; ++i) {
        cur = abs(x[i] - y[0]);
        for (j = 1; j < *y_len; ++j) {
            if (abs(x[i] - y[j]) < cur)
                cur = abs(x[i] - y[j]);
            dist[i] = cur;
        }
    }
}
```

## Important details

R	C	R SEXP
logical	int *	LGLSXP
integer	int *	INTSXP
numeric	double *	REALSXP
...	...	...

- ▶ Vectors: R is 1-based, C is 0-based.
- ▶ Matrices: row-major vectors:  $x[i, j]$  in R is  $x[(i-1) + (j-1) * nrow]$  in C.
- ▶ Transient memory management
  - ▶ `Calloc`, `Free`: under complete user control – a common source of memory leaks.
  - ▶ `R_alloc`: automatically garbage-collected on return.
- ▶ Character vectors: represented as `char **`.
- ▶ Special values: `NA`, `Inf`, etc. not allowed by default (see `?C` for how to pass to C).



## The .Call interface

```
> noquote(names(as.list(args(.Call))))
```

```
[1] name      ...      PACKAGE
```

*name* Name of the C routine.

*...* Arguments to *name*, in order (names are *not* matched to C argument names).

*PACKAGE* Look for *name* in the dll of the specified package.

## .Call example I

For each element  $x[i]$ , find the *maximum* of  $x[i]-y[j]$  over all  $j$ . E.g., in file R/maximumDistance.R

```
> maximumDistance <- function(x, y) {  
+   if (!all(is.finite(x)) || !all(is.finite(y)))  
+     stop("'x', 'y' must not be NA, NaN, Inf, -Inf")  
+   .Call("max_dist", x, y)  
+ }
```

- ▶ Information about  $x$ ,  $y$  (e.g., length) accessible at the C level.
- ▶ Return value of `.Call` is the return value of `max_dist`.

## .Call example II

```
#include <R.h>
#include <Rinternals.h>

SEXP max_dist(SEXP x_sxp, SEXP y_sxp) {
  if (isReal(x_sxp) == FALSE)
    Rf_error("'x' must be 'double'");
  if (isReal(y_sxp) == FALSE)
    Rf_error("'y' must be 'double'");

  int x_len = LENGTH(x_sxp),
      y_len = LENGTH(y_sxp);

  SEXP dist_sxp;
  PROTECT(dist_sxp = allocVector(REALSXP, x_len));

  double *x = REAL(x_sxp), *y = REAL(y_sxp),
          *dist = REAL(dist_sxp);
```

## .Call example III

```
int i, j;
double cur;
for (i = 0; i < x_len; ++i) {
    cur = 0;
    for (j = 0; j < y_len; ++j) {
        if (abs(x[i] - y[j]) > cur)
            cur = abs(x[i] - y[j]);
    }
    dist[i] = cur;
}

UNPROTECT(1);
return dist_sxp;
}
```

## Details and use

- ▶ PROTECT all SEXPs created in C; a LISTSXP needs to be protected, but not its elements.
- ▶ Main interface defined in `Rinternals.h`; alternative in `Rdefines.h`.
- ▶ Additional interface (also relevant to `.C` programming) exposed in `R_ext/*.h`

### Use

```
% R CMD SHLIB src/dist
% R --vanilla
> dyn.load("src/dist")
> source("R/distance.R")
> minimumDistance(1:10, 2:5)
> dyn.unload("src/dist")
```

# Debugging C code

## Configuring R

- ▶ Linux: `CFLAGS="-g -O0" $R_SRC/configure`
- ▶ Windows: Edit `$R_SRC/src/gnuwin32/Makefile` to read `OPTFLAGS=-O0 -Wall -pedantic`, then  
`cd $R_SRC/src/gnuwin32`  
`make DEBUG=T`

## Debug

- ▶ Start R, load dynamic library, attach debugger, set break points (details are system- and debugger specific)

# Interfaces to other languages

- ▶ `.Fortran`: like `.C`.
- ▶ Other languages possible (e.g., Java, Python) via user-contributed packages (e.g., `rJava`); usually offer R-level interface to `call-` or `eval-`like functions.