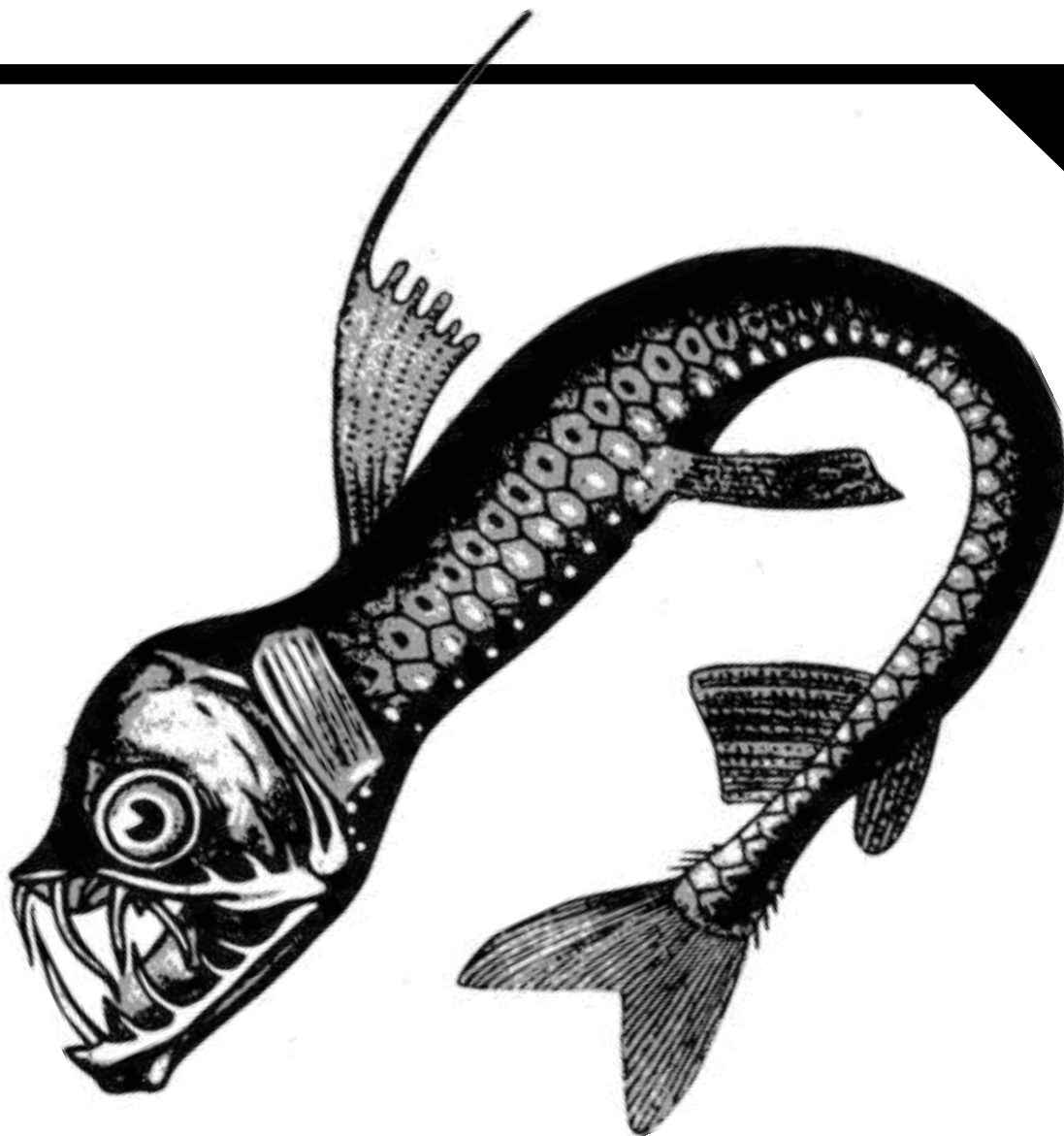


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Version  
0.1.0



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# Guide to the RNACI Package

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*Utilities for R's Native C Interface*

*Drew Schmidt and Wei-Chen Chen*

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# GUIDE TO THE RNACI PACKAGE

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UTILITIES FOR R'S NATIVE C INTERFACE

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VERSION 0.1.0

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## 1 Introduction

In this vignette, we will introduce the **RNACI** package, including its motivations and purpose, how to install it, its interface specifications, and a detailed example.

### 1.1 Purpose

This package is intended to serve somewhat the same purpose as the very (deservedly!) popular package **Rcpp**. However, this package is not meant to be a competitor to **Rcpp**. Rather, it is meant to fill a very small niche that **Rcpp** does not fill.

### 1.2 License

The **RNACI** package is licensed under the very permissive 2-clause BSD license, commonly referred to as the FreeBSD license. For a copy of the license, see the file named `LICENSE` in the root directory of the package source.

### 1.3 Installation

The package should install without issue from the command line via the usual commands:

Shell Command

```
R CMD INSTALL RNACI_0.1.0.tar.gz
```

## 2 Linking with RNACI

### 2.1 Configuring a Package to use RNACI

In your `configure.ac` and/or `src/Makevars` file(s), you can get the package linking and include information via:

```
R_SCMD="${R_HOME}/bin/Rscript -e"  
  
RNACI_LDFLAGS='${R_SCMD} "RNACI:::ldflags()" '  
RNACI_CPPFLAGS='${R_SCMD} "RNACI:::cppflags()" '
```

and adding `$(RNACI)` to `PKG_LIBS` and `$(RNACI_CPPFLAGS)` to `PKG_CPPFLAGS`. See the [Writing R Extensions](#) manual for more information. You can also see the **pbdBASE** and **pbdDMAT** packages as examples.

## 2.2 Testing the Configuration

To ensure that the package configuration is correct, you can use this test code. Include this C file:

rnaci\_test.c

```
#include <RNACI.h>

SEXP rnaci_test(SEXP mat)
{
    PRINT(mat);

    return RNULL;
}
```

and this R file:

rnaci\_test.r

```
1 rnaci_test <- function() .Call("rnaci_test", matrix(1:30, 10))
```

Then build your package with the usual R CMD INSTALL and test by loading R and running:

```
1 library(<my package>)
2
3 rnaci_test()
```

## 3 Example Usage

### 3.1 Basic Example

Suppose you have a C function `int fib(int n)` which produces the *n*'th Fibonacci number:

Fibonacci

```
int fib(n)
{
    if (n == 0 || n == 1)
        return 1;
    else
        return fib(n-1) + fib(n-1);
}
```

To wrap this for R with **RNACI**, you

Fibonacci Wrapper

```
#include <RNACI.h>
```

```
SEXP fib_wrap(SEXP n)
{
    R_INIT;
    SEXP ret;
    newRvec(ret, 1, "int");

    INT(ret) = fib(INT(n));

    R_END;
    return ret;
}
```

This may look verbose — and really it is — but for complicated examples, the tools begin to shine. So far, what we have seen isn’t really “better”, just different (although I obviously prefer it). The convenience offered by **RNACI** begins to show itself when we deal with managing lists, however.

Suppose we wanted, for example, to return both *n* and the *n*’th Fibonacci number in a list. In R’s native C interface, this is a labyrinthine nightmare, but managing return lists is very simple with **RNACI**:

#### Fibonacci Wrapper 2

```
#include <RNACI.h>

SEXP fib_wrap(SEXP n)
{
    R_INIT;
    SEXP R_list, R_list_names, nth_fib;
    newRvec(nth_fib, 1, "int");

    INT(nth_fib) = fib(INT(n));

    R_list_names = make_list_names(2, "n", "nth.fib");
    R_list = make_list(R_list_names, 2, n, nth_fib);

    R_END;
    return R_list;
}
```

You probably know that really I should be instituting a copy of *n* here; this is true of **Rcpp** and the pure native C interface as well.

Note that `INT(n)` is a shorthand for `INT(n, 0)`, which is itself shorthand for R’s `INTEGER(n)[0]`. Other values may be substituted, e.g. `INT(n, i)` for `INTEGER(n)[i]`. See [Section 4](#) for more information.

A native version of this simple example (without **RNACI**) might look like:

#### Fibonacci Wrapper 2 - Native Interface

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

SEXP fib_wrap(SEXP n)
{
    SEXP R_list, R_list_names, nth_fib;
    PROTECT(nth_fib = allocVector(INTSXP, 1));

    INTEGER(nth_fib)[0] = fib(INTEGER(n)[0]);

    PROTECT(R_list = allocVector(VECSXP, 2));
    PROTECT(R_list_names = allocVector(STRSXP, 2));

    SET_VECTOR_ELT(R_list, 0, n);
    SET_VECTOR_ELT(R_list, 1, nth_fib);

    SET_STRING_ELT(RET_NAMES, 0, mkChar("n"));
    SET_STRING_ELT(RET_NAMES, 1, mkChar("nth.fib"));

    UNPROTECT(3)
    return R_list;
}
```

## 4 Specification

If you are familiar with R's native C interface already, then you may simply wish to view `src/RNACI.h` of the **RNACI** source tree.

### 4.1 GC and Allocation

Instead of explicitly calling `UNPROTECT` on the number of `PROTECT`'d objects, start every `SEXP` function with `R_INIT` and end every `SEXP` function with `R_END`:

#### GC Counter

```
R_INIT; // Initialize the GC counter

R_END; // Call UNPROTECT on the appropriate number
```

Additionally, you don't need to call `PROTECT` anymore on new allocations; simply call `newR*` for the type of R object you want to construct:



## Allocation

```

SEXP x;
int m, n;

// Construct R list object
newRlist(x, n); // RNACI
PROTECT(x = allocVector(VECSXP, n)); // Native equiv.

// Construct numeric R vector with C-type 'type'
newRvec(x, n, type); // RNACI
PROTECT(x = allocVector(<SEXPTYPE>, n)); // Native equiv.

// Construct numeric R matrix
newRmat(x, m, n, type); // RNACI
PROTECT(x = allocMatrix(<SEXPTYPE>, m, n)); // Native equiv.

```

## Allocation Examples

```

// A length 10 integer vector
newRvec(x, 10, "int"); // RNACI
PROTECT(x = allocVector(INTSXP, 10)); // Native equiv.

// A 5 by 2 integer matrix
newRmat(x, 5, 2, "int"); // RNACI
PROTECT(x = allocMatrix(INTSXP, 5, 2)); // Native equiv.

```

If you need to protect something you aren't allocating, use `PT()` instead of `PROTECT()`; this will automatically increment the counter which keeps track of the number of `PROTECT`'d objects.

## 4.2 Accessing SEXP Data

### RNACI Data Accessors

```
SEXP x;
int i, j;

// Pointer to data
INTP(x)
DBLP(x)

// Vector data
INT(x)
INT(x, i)
DBL(x)
DBL(x, i)
STR(x)
STR(x, i)

// Matrix data
MatINT(x, i, j)
MatDBL(x, i, j)
```

### Native Data Accessors

```
SEXP x;
int i, j;

// Pointer to data
INTEGER(x)
REAL(x)

// Vector Data
INTEGER(x)[0]
INTEGER(x)[i]
REAL(x)[0]
REAL(x)[i]
(char*)CHAR(STRING_ELT(x,0))
(char*)CHAR(STRING_ELT(x,i))

// Matrix data
INTEGER(x)[i+nrows(x)*j]
REAL(x)[i+nrows(x)*j]
```

## 4.3 Lists and Dataframes

For this section, only the **RNACI** version is provided, because the native equivalent is too horrifying to mention.

### Lists

```
SEXP R_list, R_list_names;
const int num.items = 3;

// pretended we creates SEXP's item1, item2, and item3

R_list_names = make_list_names(num.items, "name1", "name2",
                                "name3"); // as many as you like
R_list = make_list(R_list_names, num.items, item1, item2, item3);
```

This dataframe example uses R default row and column names. You can probably fill in the details from the above for the named case.

### Dataframes

```
SEXP R_df;
const int num.cols = 2;
```

```
// pretended we creates SEXP's col1 and col2

R_df = make_dataframe(RNULL, RNULL, num.names, col1, col2);
```

## 4.4 Utility Functions

### Testers

```
int is_Rnull(SEXP x);
int is_Rnan(SEXP x);
int is_Rna(SEXP x);
int is_double(SEXP x);
int is_integer(SEXP x);
```

### Misc

```
RNULL // RNACI
R_NilValue // Native equiv.

// Print any SEXP with R's printing
SEXP x;
PRINT(x)
```

## 5 Q&A

### 5.1 Why make this?

Probably my biggest motivator was fun; I wanted to. Another, more pragmatic reason is that part of my workload prevents me from using **Rcpp**. I deal a lot with C and Fortran; trying to integrate C++ into that mess is not fun, and so for me, **Rcpp** is more of a burden than a savior. This leaves me stuck with the native C interface for R. And I don't like R's native C interface. This package is my attempt to make that interface (slightly) more friendly and convenient to work with.

### 5.2 Native C Interface?

Every R object is an **SEXP** (short for S-expression) object, which is a C-level struct pointer. This is explained in the [R Internals](#) manual. This package is a collection of tools for more easily managing SEXP objects.

### 5.3 Is this now, or will this ever be a competitor to **Rcpp**?

In terms of features, no. **Rcpp** is very good at aiding the writing of new code, or translating R code into compiled code. That is not a goal of this package.

### 5.4 How does this differ from **Rcpp**?

Each of these packages makes an attempt at solving a serious problem with utilizing compiled code from R: the native interface for C code in R sucks. There are huge differences between the two packages, however. In short, **Rcpp** is *much* a much more comprehensive solution. If you are new to using compiled code with R, frankly this package probably is not for you; you would likely be much better served by **Rcpp**.

**RNACI** is primarily meant for those who are working in pure C, writing C/Fortran code and wrapping it back into R. This makes the wrapping a bit simpler. It does not make the translation of R code into compiled code any simpler, which is one of **Rcpp**'s main goals.

**RNACI** is a pure C solution. **Rcpp** is a C++ solution. In the author's opinion, C++ brings a lot of needless complications and headaches if you aren't actually using C++; that is, if you are only writing C and Fortran, then bringing in C++ **will** create problems.

Another important difference is that **RNACI** is more permissively licensed than **Rcpp** (BSD rather than GPL). If this issue is important to you but you live in the C++ world, you may be interested in Romain Francois' new **Rcpp11** package.

### 5.5 Why would I want to use this package?

If spend a lot of time bringing code to R (especially if you deal with lists and dataframes), this package offers a lot of convenient shorthand for doing so.

### 5.6 Is this really easier than R's native interface?

It is for me, no question; notably, returning lists and dataframes is *much* less painful through some [stdarg](#) sorcery. Most of the rest of the package is minor cosmetic things; but the package is very permissively licensed, so feel free to pick and choose what you want, however you want.

### 5.7 How would I use **RNACI** in a package?

Assuming that you have some compiled code you have or want to create to use with a package, you simply link with **RNACI** and then wrap that compiled code with the utilities provided by **RNACI**. For the former, see [Section 2](#), and for the latter, see [Section 4](#) and [Section 3](#). For actual package examples using **RNACI**, see [pbdBASE](#) version  $\geq 0.3-0$ , [pbdDMAT](#) version  $\geq 0.3-0$ , and [memuse](#) version  $\geq 2.0$ .

---

Philosophically, you should never have the bulk of the work of a function (of any importance) be handled by the `R` interface (including **RNACI**'s version of it). If you do, then your code can never (easily) have a life outside of `R`. That may sound fine to you now, but if you ever decide to take some of your work outside of `R`, then you can't easily take your compiled code with you. This is just bad practice and shortsightedness.