

Using Rcpp* packages for easy and fast extension of R with C++



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Outline

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- Rcpp : extending R with C++
- RcppGSL for fast random draws
- RcppArmadillo for high-performance linear algebra

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4 Build of R packages using Rcpp

- Useful functions
- Example of package built with Rcpp



Rcpp R package

- **Rcpp** is an R package to extend R with C++ code
- Main advantage : C++ is fast, it accelerates R (see next sections)
- Written by **Dirk EDDELBUETTEL** and **Romain FRANCOIS**
- <http://www.rcpp.org/>



Simple Rcpp example

C++ code (in file Code/addition.cpp)

```
#include <Rcpp.h>
using namespace Rcpp;
// [[Rcpp::export]]
int addition(int a, int b) {
  return a + b;
}
```

R code

```
Rcpp::sourceCpp("Code/addition.cpp")
addition(2, 2)
```

```
## [1] 4
```



Rcpp advantages

Thanks to `Rcpp::sourceCpp()`

- Compile the C++ code
- Export the function to the R session
- Direct interchange of R objects (including S3, S4) between R and C++
- ... (many more, see `vignette("Rcpp-package")`)



GSL and RcppGSL



GNU Scientific Library

- Numerical library for C and C++ programmers
- Reliable random number generator algorithms
- Thoroughly tested and fast random number distributions
- Linear algebra (matrices and vectors)
- <https://www.gnu.org/software/gsl/>

RcppGSL

- Interface between R and GSL
- Using Rcpp to interface R and C
- <http://dirk.eddelbuettel.com/code/rcpp.gsl.html>



GSL random number distributions

- GSL v2.6 includes **38 random number distributions** (see [GNU GSL](#))
- It's easy to implement additional random number distributions from the GSL base distributions (e.g. truncated normal distribution)
- For comparison, R API includes "only" 24 random number distributions (see [Writing R Extensions](#))
- Random draws are faster with GSL than with R (eg. `gsl_ran_gamma()` vs. `R::rgamma()`)



RcppGSL example

C++ code

```
#include <Rcpp.h>
#include <gsl/gsl_rng.h>
#include <gsl/gsl_randist.h>
using namespace Rcpp;
// [[Rcpp::depends(RcppGSL)]]
// [[Rcpp::export]]
Rcpp::NumericVector my_rnorm(int nsamp, double mu,
                             double sigma) {
  gsl_rng *s = gsl_rng_alloc(gsl_rng_mt19937); // Random seed
  Rcpp::NumericVector beta(nsamp);
  for (int i = 0; i < nsamp; i++) {
    beta[i] = mu + gsl_ran_gaussian(s, sigma); // Random draw
  }
  return beta;
}
```



RcppGSL example

R code

```
library(Rcpp)
library(RcppGSL)
beta <- my_rnorm(100, 5, 2)
par(cex=2)
hist(beta)
```



Armadillo and RcppArmadillo

Armadillo

- C++ library for linear algebra and scientific computing
- Provides high-level syntax and functionality : speed and ease of use
- Classes for vectors, matrices and cubes
- Matrix operations, matrix decomposition, linear model solver, etc.
- <http://arma.sourceforge.net/>



RcppArmadillo

- Interface between R and Armadillo
- Using Rcpp to interface R and C++
- <http://dirk.eddelbuettel.com/code/rcpp.armadillo.html>



RcppArmadillo example

C++ code

```
#include <RcppArmadillo.h>
// [[Rcpp::depends(RcppArmadillo)]]
// [[Rcpp::export]]
Rcpp::List fastLm(const arma::mat& X, const arma::colvec& y) {
    int n = X.n_rows, k = X.n_cols;

    arma::colvec coef = arma::solve(X, y);      // fit model  $y \sim X$ 
    arma::colvec res  = y - X*coef;                // residuals
    // std.errors of coefficients
    double s2 = std::inner_product(res.begin(),
                                    res.end(),
                                    res.begin(), 0.0)/(n - k);

    arma::colvec std_err = arma::sqrt(s2 *
        arma::diagvec(arma::pinv(arma::trans(X)*X)));
    return Rcpp::List::create(Rcpp::Named("coefficients") = coef,
                            Rcpp::Named("stderr")       = std_err,
                            Rcpp::Named("df.residual") = n - k);
}
```



RcppArmadillo example

R code

```
library(Rcpp)
library(RcppArmadillo)
# Trees data-set
y <- log(trees$Volume)
X <- cbind(1, log(trees$Girth))
# fastLm
mod <- fastLm(X, y)
mod$coef

##           [,1]
## [1,] -2.353325
## [2,]  2.199970
```



Licenses

- Licenses : GNU General Public License, Apache License 2.0 for Armadillo
- Free software licenses : we can use, modify and redistribute these softwares



Distance computation

C++ code (in file Code/arma_distmat.cpp)

```
#include <RcppArmadillo.h>
// [[Rcpp::depends(RcppArmadillo)]]
using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
arma::mat arma_distmat(const arma::mat& x) {

    int np = x.n_rows;

    arma::mat distmat; distmat.zeros(np, np);
    for (int i = 0; i < np; i++) {
        arma::vec p0 = x.row(i).t();
        for (int j = i + 1; j < np; j++) {
            arma::vec p1 = x.row(j).t();
            arma::vec diff = p0 - p1;
            double squared_diff = as_scalar(diff.t() * diff);
            distmat(j, i) = distmat(i, j) = sqrt(squared_diff);
        }
    }
    return distmat;
}
```



Distance computation

R code

```
Rcpp::sourceCpp("Code/arma_distmat.cpp")
library(rdist)
X <- matrix(rnorm(10000), ncol=2)

# Benchmark
library(rbenchmark)
Benchmark <- benchmark(
  "arma_distmat" = {arma_distmat(X)},
  "cdist" = {cdist(X,X)},
  replications=10,
  columns = c("test", "elapsed", "relative"))
knitr::kable(Benchmark, booktabs=TRUE) %>%
  kableExtra::kable_styling(latex_options=c("HOLD_position","striped"),
                            full_width=FALSE)
```

test	elapsed	relative
arma_distmat	5.788	1.00
cdist	15.104	2.61

Simple linear regression

Short presentation of Rcpp* packages

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Application examples

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Exercise to do yourself

Build of R packages using Rcpp

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Functions to build an R package

- `Rcpp.package.skeleton()` to generate a new Rcpp package (modifying `DESCRIPTION` and `NAMESPACE`)
- `Rcpp::compileAttributes()` scans the C++ files and generates the `RcppExports.cpp` file to make the functions preceded by `// [[Rcpp::export]]` available in R.



jSDM R package

jSDM 0.1.0 Get started Reference Articles ▾ Change log

jSDM R Package

Package for fitting joint species distribution models (jSDM) in a hierarchical Bayesian framework (Warton *et al.* 2015). The Gibbs sampler is written in C++. It uses Rcpp, Armadillo and GSL to maximize computation efficiency.



Links

Download from CRAN at

<https://cloud.r-project.org/package=jSDM>

Browse source code at

<https://github.com/ghislainv/jSDM>

Report a bug at

<https://github.com/ghislainv/jSDM/issues>

License

GPL-3 | file LICENSE

Developers

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Copyright holder, funder

Dev status

build passing

CRAN 0.1.0

DOI 10.5281/zenodo.3253466

do wnloads 231/month

System requirements

Make sure the GNU Scientific Library (GSL) is installed on your system.

Installation

Install the latest stable version of jSDM from CRAN with:

```
install.packages("jSDM")
```

Or install the development version of jSDM from GitHub with:

```
devtools::install_github("ghislainv/jSDM")
```

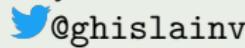
References

Warton, D.I., Blanchet, F.G., O'Hara, R.B., Ovaskainen, O., Taskinen, S., Walker, S.C. & Hui, F.K. (2015) So many variables: Joint modeling in community ecology. *Trends in Ecology & Evolution*, **30**, 766–779.

- <https://ecology.ghislainv.fr/jSDM>
- Made with Rcpp* packages



... Thank you for attention ...



<https://ecology.ghislainv.fr>

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