

# PARALLEL R

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- ▶ It is a package originally develop by Dirk Eddelbuettel and Romain François
- ▶ It aims to ease the extension of R with C++ code.
- ▶ It allows to load C++ code in an interactive session.
- ▶ It has framework to help when creating package with Rcpp.

- ▶ You can create function directly in R code as a string put into `cppFunction()` call.
  - ▶ In this case Rcpp will do most of the heavy lifting for You. (headers, compilation, linking)
- ▶ You can write C++ function and source it by calling `sourceCpp()`.
  - ▶ Adding `verbose = TRUE` will show the whole process.
- ▶ Creating a package with C++ files with `// [[Rcpp::export]]` attribute.

- ▶ Rcpp syntactic sugar makes rewrite from R code easier, thanks to possibility to sometimes use R like notation.
- ▶ Rcpp attributes allows not just easy C++ function exports, but also define dependencies, change function names, define initialization functions.
- ▶ Rcpp contains function to handle exception in the C++ code and to check for user interruption.
- ▶ There is whole ecosystem of Rcpp packages that further extends its capabilities.

- ▶ A Brief Introduction to Rcpp
- ▶ Rcpp Attributes
- ▶ Writing a package that uses Rcpp
- ▶ Rcpp syntactic sugar

To use the Rcpp package in a C++ file and export a C++ function to R, you need to follow these steps:

- ▶ Install the Rcpp package if it is not already installed. You can do this from within R by running `install.packages("Rcpp")`.
- ▶ Create a new `.cpp` file with your C++ code.
- ▶ Add `Rcpp::export` attribute to functions that should be available for use from R.

- ▶ For CUDA we need to add specific `.cu` file containing the code for the GPU. Such functions are called kernel functions.
- ▶ To be able to call this kernel function in our C++ function, we will need to create `.h` file called header file, which describes the function we want to call.
- ▶ Additionally, it is necessary to configure `makevars` file which tells R which compiler to use

- ▶ R function that calls a C++ function.
- ▶ C++ file
- ▶ cu file
- ▶ makevars
- ▶ header file
- ▶ DESCRIPTION
- ▶ NAMESPACE



Mandelbrot set is defined as  $f_c(z) = z^2 + c$ , where  $c$  is a complex number corresponding to the point coordinates.

```
mandelbrot <- function(c, max_iter = 100) {  
  z <- c  
  for (i in 1:max_iter - 1) {  
    z <- z ^ 2 + c  
    if (abs(z) > 2) {  
      return(i)  
    }  
  }  
  return(max_iter)  
}
```

```
dc <- cmax - cmin
x <- y <- 1:resolution - 1
x <- Re(cmin) + (x / resolution * Re(dc))
y <- Im(cmin) + (y / resolution * Im(dc))
points <-
  outer(x, y, function(x, y)
    complex(real = x, imaginary = y))
result_for <- matrix(NA,
                     dim(points)[1], dim(points)[2])
for (x in 1:dim(points)[1]) {
  for (y in 1:dim(points)[2]) {
    result_for[x, y] <- mandelbrot(points[x, y],
                                   max_iter)
  }
}
```

```
int Mandel(double real, double im,
           int max_iter = 100)
{
    std::complex<double> c(real, im);
    std::complex<double> z = c;
    for (int i=0; i< max_iter; i++){
        z = z * z + c;
        if (std::abs(z) > 2) {
            return i;
        }
    }
    return max_iter;
}
```

```
...  
std::complex<double> dc = cmax - cmin;  
IntegerMatrix out( resolution );  
for (int i=0; i < resolution; i++){  
    for(int j=0; j < resolution; j++){  
        double helper = static_cast<double>(i);  
        double helper2 = static_cast<double>(j);  
        double fx = helper / resolution * real(dc);  
        double fy = helper2 / resolution * imag(dc);  
        std::complex<double> c(real(cmin) + fx,  
                                imag(cmin) + fy);  
        ...  
    }  
}  
...
```