1 Preliminaries

Some key things that need to be installed are: *Xcode* or at least *command line tools* (less memory requirements). For the latter, you can register as an Apple developer for free at `https://developer.apple.com` to download it. Note that the latest version of *Xcode* might not be compatible with your OS. *Homebrew*, and then *GSL* tools via command line in the following way

```
brew install gsl
brew link gsl
```

If there is a homebrew error about the directory being unwritable use `brew doctor` first to confirm that's the problem followed by

```
sudo chown -R maria:admin /usr/local/lib/pkgconfig
```

to make the directory writable. Then, in the RStudio prompt

```
install.packages('gsl',type = 'source')
install.packages(Rcpp)
install.packages(RcppGSL)
install.packages(inline)
```

2 Two possible ways of building a wrap for C++ code

1. Inline package , writing a bit of simple C++ code and then wrapping it with `cxxfunction`

   ```
   require(inline)
   incltxt<-'
   int fibonacci(const int x){
   if (x == 0) return(0);
   if (x == 1) return(1);
   return fibonacci(x-1)+fibonacci(x-2);
   }'

   fibRcpp <- cxxfunction(signature(xs="int"), plugin="Rcpp",
   incl = incltxt, body = 'int x = Rcpp::as<int>(xs);
                           return Rcpp::wrap(fibonacci(x) );')
    # For evaluating the function, say we want the element number 10
    # of the Fibonacci sequence:
    fibRcpp(10)
   ```

2. Using `//[[Rcpp::export]]`, also called the Rcpp attributes.

   ```
   # includee<Rcpp.h>
   using namespace Rcpp;

   //[[Rcpp::export]]
   int fibonacci(const int x){
   if (x<2)
   return x;
   else
   return (fibonacci(x-1)+fibonacci(x-2));
   }
   ```

and then just use `sourceCpp("fibonacci.cpp")` to source it directly. Another example of the `inline` way, where we set an integer vector pointer where you specify each of the entries

```
src<-'
Rccp::IntegerVector epn(4);
epn[0] = 6;
epn[1] = 14;
epn[2] = 496;
epn[3] =8182;
return epn;'
fun<-cxxfunction(signature(),src,plugin = "Rcpp")
```

Other data types are:

```
Rcpp::IntegerVector
Rcpp::NumericVector
Rcpp::LogicalVector
Rcpp::CharacterVector
Rcpp::ExpressionVector
Rcpp::RawVector
Rcpp::IntegerMatrix
Rcpp::NumericMatrix
```

This is another example of different ways to build pointers

```
src<-'
Rcpp::NumericVector vec(vx);
double p = Rcpp::as<double>(dd);
double sum = 0.0;
for (int i =0; i<vec.size();i++) {
sum +=pow(vec[i],p);
}
return Rcpp::wrap(sum);'
fun<-cxxfunction(signature(vx="numeric",))
```

Note that, because of C++ funny things, you can't use the same name for an input and an output. The `clone` command is used to overcome this problem.

```
src<-'
Rcpp::NumericVector invec(vx);
Rcpp::NumericVector outvec = Rcpp::clone(vx);
for(i=0: i<invec.size(); i++>){
outvec[i] = log(invec[i]);
}
return outvec;'
fun <-cxxfunction(signature(vx="numeric"),src,plugin="Rcpp")
return outvec
```

⋮

Add something about the list class for general purpose pointers

⋮

Note that all of the above are very simple C++ bits of code with no dependencies to `#includes` `studio.h` libraries. It looks like once these dependencies need to be used then it won't work as in *Python* or *Matlab* but a `Rcpp.package.skeleton()` needs to be created and all *.h* libraries should be saved in the source folder in order to be able to compile the *.cpp* files. This point will be discussed in the next section.

3   Build a package that contains Rcpp files

A skeleton package is a minimal package providing a working example which can then be adapted and extended as needed by the user.

```
#To create the package
require(Rcpp)
Rcpp.package.skeleton("RGLFM")
# To compile it
compileAttributes("~/FAP_Rpackage/GLFM/src/Ccode/wrapper_R/RGLFM",verbose=TRUE)
install.packages("~/FAP_Rpackage/GLFM/src/Ccode/wrapper_R/RGLFM", repos = NULL, type="source")
```

Manually add all .cpp and .h files to the package's `src` folder. And then you can see the wrapper in the `R` folder. To build the package and create the `.tar` file, from the command line:

```
 R CMD build RGLFM
 # Quit R here
 R CMD INSTALL RGLFM_1.0.tar.gz
  R CMD check RGLFM_1.0.tar.gz
```

then, in RStudio:

```
require(Rcpp)
setwd("~/Documents/Working_papers/FAP_Rpackage/GLFM/src/Ccode/wrapper_R/RGLFM")
library('RGLFM')
rcpp_hello_world()
```