

# Sharing early experiences of programming with Rcpp

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# Rcpp package

**Rcpp** package facilitates the application of cpp for computations in R.

It provides interface for communication between R and cpp simplifying it.

It is much easier to benefit from the best of the two worlds:

**cpp** code is compiled facilitating quick computations:  
perfect for writing functions

**R** code is interpreted and dynamic:  
perfect for data analysis using functions written in cpp

# Rcpp package

In this presentation we are focusing on applications to Bayesian estimation that relies on:

**elements of programming:** functions, loops, etc.

**compatible object types:** scalars, vectors, matrices, lists, etc.

**linear algebra:** using library Armadillo

**random number generators:** fast and reproducible using Rcpp

**Rcpp** is an R package providing the interface, object type compatibility, ..., and **vectorised random number generators**

**Armadillo** is a cpp library for linear algebra with fantastic documentation online

**RcppArmadillo** is an R package providing simplified interface with Armadillo and providing compatibility with its object types

# Rcpp package: learning resources

**book:** Eddelbuettel (2013) Seamless R and C++ Integration with Rcpp

**bookdown:** Tsuda (2020) Rcpp for everyone

**vignettes:** for packages Rcpp and RcppArmadillo  
(published in JSS, CSDA, and TAS)

**datacamp course:** Optimizing R Code with Rcpp by Romain François

**online resources:** RcppGallery, Armadillo library  
documentation, [stackoverflow.com](https://stackoverflow.com)

A simple example

# Developing a function in a .cpp file

## nicetry.cpp

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

using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
vec nicetry (mat y, mat x) {
  vec beta_hat = solve(x.t()*x, x.t()*y);
  return beta_hat;
}

/** R
x = cbind(rep(1,5),1:5)
y = x %*% c(1,2) + rnorm(5)
nicetry(y, x)
solve(crossprod(x), crossprod(x,y))
*/
```

# Using a function from a .cpp file

## nicetry.cpp

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

using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
vec nicetry (mat y, mat x) {
  vec beta_hat = solve(x.t()*x, x.t()*y);
  return beta_hat;
}
```

## linreg.R

```
library(Rcpp)
sourceCpp("nicetry.cpp")

x      = cbind(rep(1,5), 1:5)
y      = x %*% c(1,2) + rnorm(5)
beta.hat = nicetry(y, x)
```

Simulation Smoother



# Simulation smoother

Sample random draws from the multivariate normal distribution:

$$\mathcal{N}_T(\Omega^{-1}\alpha, \Omega^{-1})$$

$\Omega$  -  $T \times T$  precision matrix that is band- or tri-diagonal

$\alpha$  -  $T \times 1$  location vector

A random draw is computed by

$$L^{-1'}(L^{-1}\alpha + \epsilon)$$

$L = \mathbf{chol}(\Omega)$  - is lower-triangular

$\epsilon$  -  $T$ -vector with independent standard normal draws

# R: simulation smoother #1

rmvnorm\_solve.R

```
rmvnorm_solve = function(n, precision, location){  
  T          = dim(precision)[1]  
  precision.chol.inv = solve(t(chol(precision)))  
  
  epsilon    = matrix(rnorm(n*T), ncol=n)  
  draw       = t(precision.chol.inv) %*%  
               (matrix(rep(precision.chol.inv%*%  
                           location,n), ncol=n) + epsilon)  
  
  return(draw)  
}
```

## R: simulation smoother #2

rmvnorm\_bandchol.R

```
library(mgcv)

rmvnorm_bandchol = function(n, precision, location){
  T          = dim(precision)[1]
  precision.L = t(bandchol(precision))

  epsilon    = matrix(rnorm(n*T), ncol=n)
  a          = forwardsolve(precision.L, location)
  draw       = backsolve(t(precision.L),
                        matrix(rep(a,n), ncol=n) + epsilon)

  return(draw)
}
```

# R: simulation smoother #3

rmvnorm\_trichol.R

```
library(mgcv)

rmvnorm_trichol = function(n, precision, location){
  T          = dim(precision)[1]
  lead.diag  = diag(precision)
  sub.diag   = sdiag(precision, -1)

  precision.chol = trichol(ld = lead.diag, sd=sub.diag)
  precision.L    = diag(precision.chol$ld)
  sdiag(precision.L, -1) = precision.chol$sd

  epsilon     = matrix(rnorm(n*T), ncol=n)
  a           = forwardsolve(precision.L, location)
  draw        = backsolve(t(precision.L),
                          matrix(rep(a,n), ncol=n)
                          + epsilon)

  return(draw)
}
```

## rmvnorm\_arma\_inv.cpp

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

using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
mat rmvnorm_arma_inv (int n, mat precision, vec location){
  int T = precision.n_rows;

  mat epsilon(T, n);
  for (int i=0; i<n; i++){
    epsilon.col(i) = as<vec>(rnorm(T));
  }

  mat location_matrix(T, n, fill::zeros);
  location_matrix.each_col() += location;
  mat precision_chol_inv = trans(inv(trimatu(chol(precision))));
  mat draw = trans(precision_chol_inv) *
              (precision_chol_inv * location_matrix
               + epsilon);

  return draw;
}
```

## rmvnorm\_arma\_solve.cpp

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

using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
mat rmvnorm_arma_solve(int n, mat precision, vec location){
  int T = precision.n_rows;

  mat epsilon(T, n);
  for (int i=0; i<n; i++){
    epsilon.col(i) = as<vec>(rnorm(T));
  }

  mat location_matrix(T, n);
  location_matrix.each_col() = location;
  mat precision_chol = trimatu(arma::chol(precision));
  mat draw           = solve(precision_chol,
                             solve(trans(precision_chol),
                                      location_matrix) + epsilon);

  return draw;
}
```



# Simulation smoother

## SimulationSmoother.R

```
library(mgcv)
library(Rcpp)
library(microbenchmark)

source("rmvnorm_trichol.R")
source("rmvnorm_bandchol.R")
source("rmvnorm_solve.R")
sourceCpp("rmvnorm_arma_inv.cpp")
sourceCpp("rmvnorm_arma_solve.cpp")
sourceCpp("rmvnorm_arma_stochvol.cpp")

set.seed(12345)
n      = 100
T      = 250
s      = rgamma(1, shape=10, scale=10)
precision = rgamma(1, shape=10, scale=10)*diag(T) + 2*s*diag(T)
sdiag(precision, 1) = -s
sdiag(precision, -1) = -s
location = as.matrix(rnorm(T))
```



# Simulation smoother

## SimulationSmoother.R

```
microbenchmark(  
  R.solve      = rmvnorm_solve(n, precision, location),  
  R.band       = rmvnorm_bandchol(n, precision, location),  
  R.tridiag    = rmvnorm_trichol(n, precision, location),  
  cpp.inv      = rmvnorm_arma_inv(n, precision, location),  
  cpp.sol      = rmvnorm_arma_solve(n, precision, location),  
  cpp.sto      = rmvnorm_arma_stochvol(n, precision, location),  
  check        = "equal",  
  setup        = set.seed(123)  
)
```

Unit: milliseconds

	expr	min	lq	mean	median	uq	max	neval
	R.solve	15.012696	16.022046	19.049814	16.741754	21.296232	41.665376	100
	R.band	5.521327	6.298791	13.280026	6.719498	9.619501	117.470313	100
	R.tridiag	3.674740	4.004751	8.079238	4.449233	4.916731	178.783048	100
	cpp.inv	10.622570	11.113882	12.171498	11.594380	12.052189	21.046931	100
	cpp.sol	2.024739	2.183631	2.789788	2.808968	2.971318	6.256791	100
	cpp.sto	1.556249	1.621727	1.875903	1.882130	1.988038	3.034821	100

Gibbs sampler for a local-level model

# Gibbs sampler for a local-level model

The model for a unit-root non-stationary variable:

$$y_t = \mu_t + \epsilon_t \quad \epsilon_t \sim \mathcal{N}(0, \sigma^2)$$

$$\mu_t = \mu_{t-1} + m_t \quad m_t \sim \mathcal{N}(0, \sigma_m^2)$$

Gibbs sampler:

$$\mu|y, \mu_0, \sigma^2, \sigma_m^2 \sim \mathcal{N}_T(\overline{V}^{-1}\overline{\mu}, \overline{V}^{-1})$$

$$\overline{V}^{-1} = \sigma^{-2}I_T + \sigma_m^{-2}H'H$$

$$\overline{\mu} = \sigma^{-2}y + \sigma_m^{-2}\mu_0 e_1$$

$$\mu_0|y, \mu_1, \sigma_m^2 \sim \mathcal{N}\left((\sigma_m^{-2} + \underline{v}^{-1})^{-1}, (\sigma_m^{-2} + \underline{v}^{-1})^{-1} \sigma_m^{-2} \mu_1\right)$$

$$\sigma^2|y, \mu \sim \text{IG2}(\underline{s} + (y - \mu)'(y - \mu), \underline{v} + T)$$

$$\sigma_m^2|\mu, \mu_0 \sim \text{IG2}(\underline{s} + (H\mu - \mu_0 e_1)'(H\mu - \mu_0 e_1), \underline{v} + T)$$

## LL\_arma\_solve.cpp - part 1

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

using namespace Rcpp;
using namespace arma;

// [[Rcpp::export]]
List LL_arma_solve(
    vec y,
    int S = 10,
    Nullable<List> starting_values = R_NilValue,
    NumericVector Hyper = NumericVector::create(10,1,3)) {

    vec hyper = as<vec>(Hyper);
    int T = y.n_rows;

    vec aux_mu(T, fill::zeros);
    double aux_mu0 = 0;
    double aux_sigma2 = 1;
    double aux_sigma2m = 1;
```

## LL\_arma\_solve.cpp - part 2

```
if (starting_values.isNotNull()){
  List Starting_values(starting_values);
  aux_mu           = as<vec>(Starting_values["mu"]);
  aux_mu0          = Starting_values["mu0"];
  aux_sigma2       = Starting_values["sigma2"];
  aux_sigma2m      = Starting_values["sigma2"];
}

mat posterior_mu(T, S);
vec posterior_mu0(S);
vec posterior_sigma2(S);
vec posterior_sigma2m(S);

mat H(T, T, fill::eye);
H.diag(-1) += -1;
mat HH = H.t() * H;
mat IT(T, T, fill::eye);
```

## LL\_arma\_solve.cpp - part 3

```
for (int s=0; s<S; s++){
    double vm      = 1/((1/hyper[0]) + (1/aux_sigma2m));
    aux_mu0        = R::rnorm((vm*aux_mu[0])/aux_sigma2m, sqrt(vm));

    double res_ss   = sum(pow(y - aux_mu,2));
    aux_sigma2      = (hyper[1] + res_ss)/R::rchisq(hyper[2] + T);

    vec mu0_vec(1, fill::value(aux_mu0));
    double mu_ss    = sum(pow(diff(join_cols(mu0_vec, aux_mu)), 2));
    aux_sigma2m     = (hyper[1] + mu_ss)/R::rchisq(hyper[2] + T);

    mat precision   = IT/aux_sigma2 + HH/aux_sigma2m;
    vec location    = y/aux_sigma2 + aux_mu0*IT.col(0)/aux_sigma2m;
    mat precision_chol = trimatu(chol(precision));
    vec epsilon     = rnorm(T);
    aux_mu          = solve(precision_chol,
                           solve(trans(precision_chol),
                                location) + epsilon);

    posterior_mu.col(s)      = aux_mu;
    posterior_mu0(s)        = aux_mu0;
    posterior_sigma2(s)     = aux_sigma2;
    posterior_sigma2m(s)    = aux_sigma2m;
}
```

## LL\_arma\_solve.cpp - part 4

```
List last_draw;
last_draw["mu"]      = aux_mu;
last_draw["mu0"]     = aux_mu0;
last_draw["sigma2"]  = aux_sigma2;
last_draw["sigma2m"] = aux_sigma2m;

List posterior;
posterior["mu"]      = posterior_mu;
posterior["mu0"]     = posterior_mu0;
posterior["sigma2"]  = posterior_sigma2;
posterior["sigma2m"] = posterior_sigma2m;

List output;
output["last.draw"]  = last_draw;
output["posterior"]  = posterior;

return output;
}
```

# Gibbs sampler for a local-level model

LL.R

```
microbenchmark(R.not      = LL_nothing_special(y250, S),
               R.ban      = LL_band_precision(y250, S),
               R.tri      = LL_tridiag_precision(y250, S),
               cpp.inv     = LL_arma_inv(y250, S),
               cpp.sol     = LL_arma_solve(y250, S),
               cpp.sto     = LL_arma_stochvol(y250, S),
               setup      = set.seed(123)
               )
```

Unit: milliseconds

expr	min	lq	mean	median	uq	max	neval
R.not	127.266986	132.328077	155.005314	138.435266	154.022634	327.33741	100
R.ban	46.167909	49.679159	95.663136	55.453045	159.034735	324.58272	100
R.tri	19.089019	22.110003	41.142920	26.470268	29.056582	150.71913	100
cpp.inv	10.580124	11.180750	12.486657	11.937587	12.845465	26.01354	100
cpp.sol	10.542156	11.145461	12.291518	11.682092	12.380952	27.42871	100
cpp.sto	8.177979	8.542281	9.554215	9.115295	9.739227	19.19876	100



What's next?

Rewrite all your code in Rcpp!