



OPTIMIZING R CODE WITH RCPP

Random number generation

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Generating single random numbers

```
// one number from a N(0,1)
double x = R::rnorm( 0, 1 ) ;

// one number from a U(-2,2)
double y = R::runif( -2, 2 ) ;

// ...
```



Generating vectors

Random number generators in the `Rcpp::` namespace.

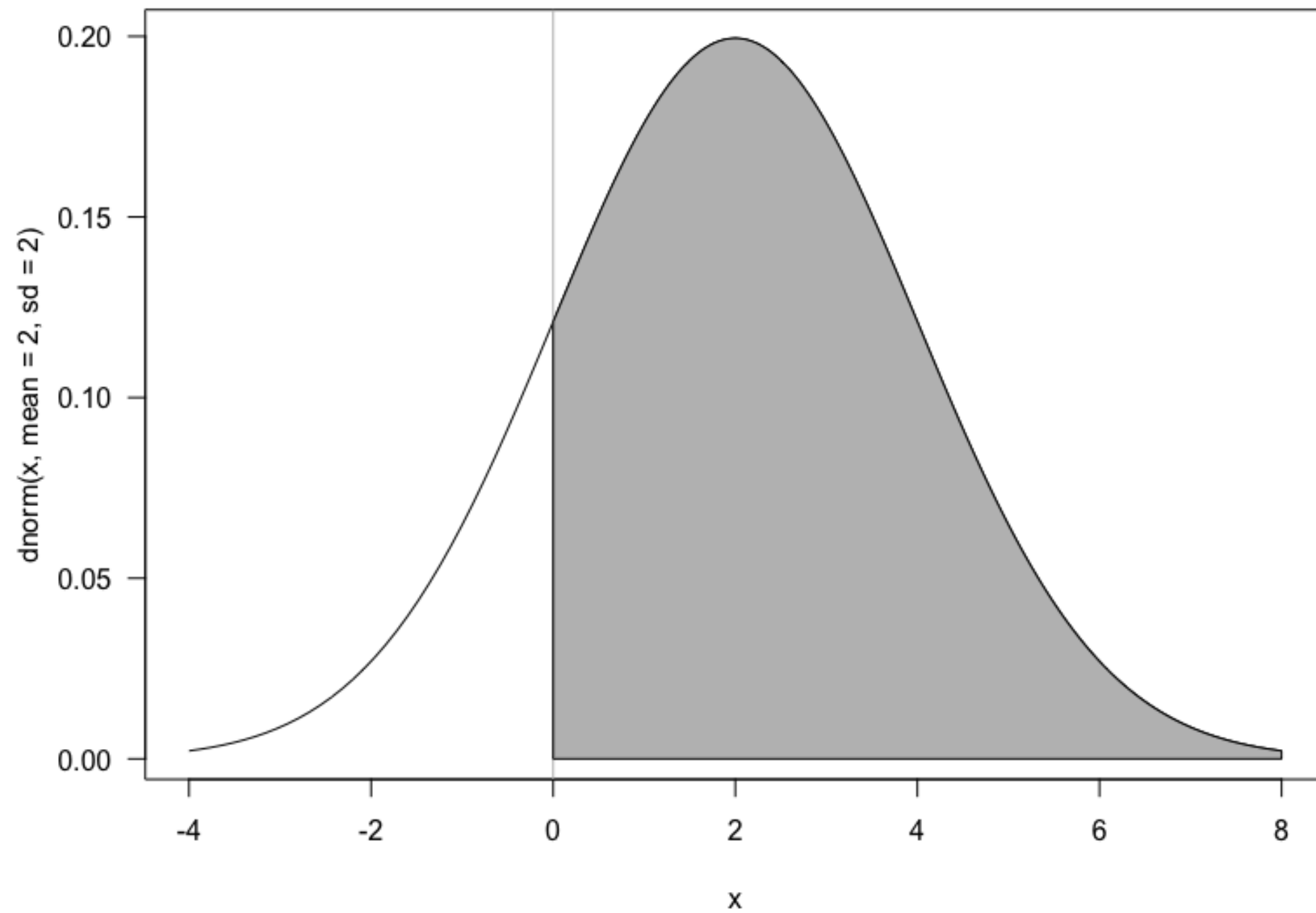
```
NumericVector x = rnorm(10, 0, 2) ;  
// same as this below  
// because of using namespace Rcpp ;  
//  
// NumericVector x = Rcpp::rnorm(10, 0, 2) ;
```

Alternative using scalar versions from `R::`

```
// same as  
NumericVector x(10) ;  
for(int i=0; i<10; i++){  
    x[i] = R::rnorm(0, 2) ;  
}
```



Truncated $N(2,2)$ at $x > 0$



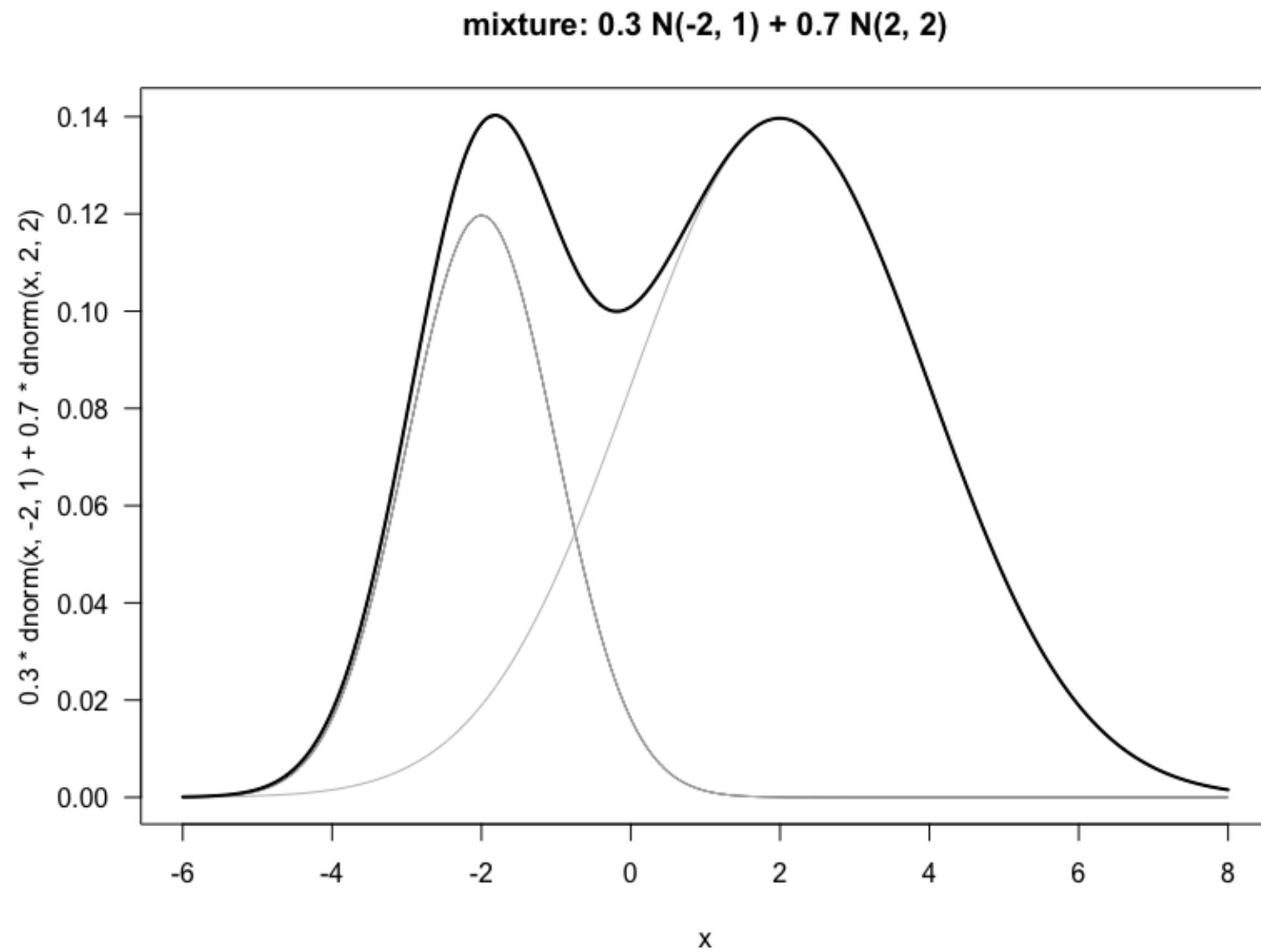


Rejection sampling

```
// we generate n numbers
NumericVector x(n) ;

// fill the vector in a loop
for( int i=0; i<n; i++){
    // keep generating d until it gets positive
    double d ;
    do {
        d = ... ;
    } while( d < 0 ) ;

    x[i] = d ;
}
```





Generate from a mixture of distributions

- Choose the component of the mixture using the weights

```
int component( NumericVector weights, double total_weight ){  
    // return the index of the selected component  
}
```

- Generate the number using the parameters of the selected components

```
NumericVector rmix( int n, NumericVector weights, NumericVector means,  
                    NumericVector sds ){  
  
    NumericVector res(n) ;  
  
    for( int i=0; i<n; i++){  
        // find which component to use  
        ...  
  
        // simulate using the mean and sd from the selected component  
        ...  
    }  
  
    return res ;  
}
```



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Let's practice!



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Rolling operations

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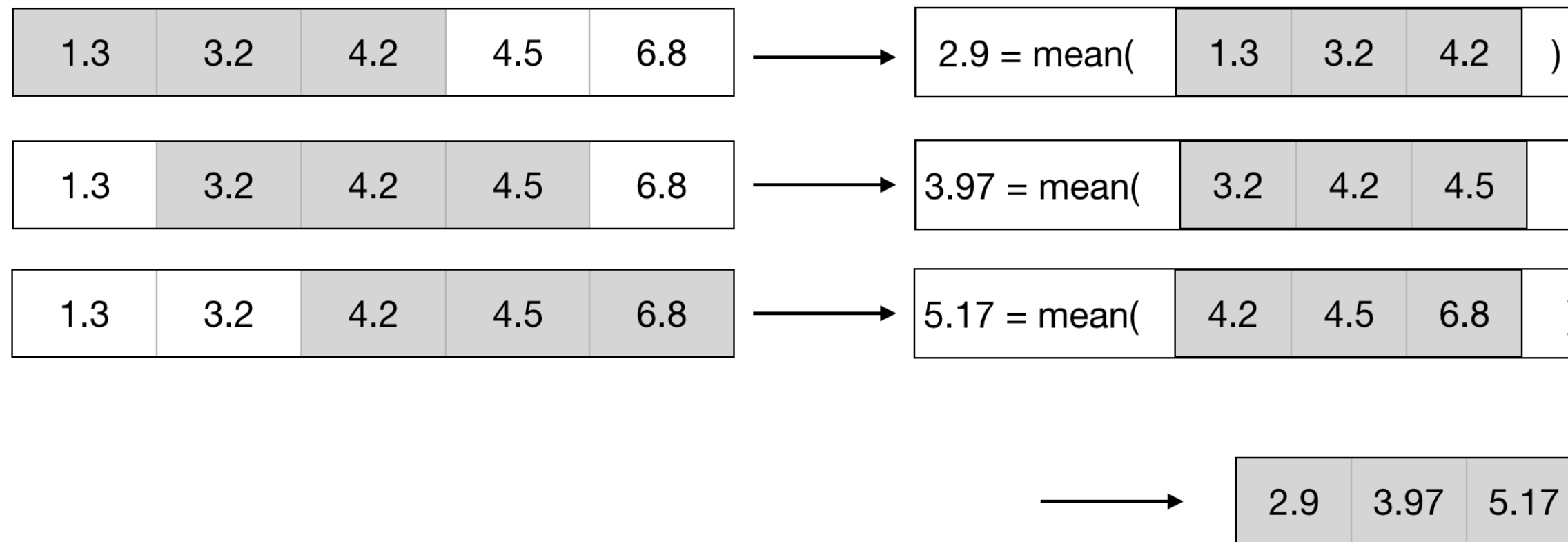


Rolling means

```
rollmean1 <- function(x, window = 3) {  
  n <- length(x)  
  
  # create empty vector full of NA  
  res <- rep(NA, n)  
  
  # fill the values  
  for( i in seq(window, n) ){  
    idx <- seq(i-window+1, i)  
    res[i] <- mean(x[idx])  
  }  
  res  
}
```



Rolling means



- Make an integer vector to hold indice, extract the relevant part of `x`
- Call the `mean` function on that extract



Alternative algorithm

1.3	3.2	4.2	4.5	6.8
-----	-----	-----	-----	-----

$$8.7 = 1.3 + 3.2 + 4.2$$

2.9

-			+	
1.3	3.2	4.2	4.5	6.8

$$11.9 = 8.7 - 1.3 + 4.5$$

3.97

	-			+
1.3	3.2	4.2	4.5	6.8

$$15.5 = 11.9 - 3.2 + 6.8$$

5.17



Alternative algorithm

```
rollmean2 <- function(x, window = 3) {  
  n <- length(x)  
  res <- rep(NA, n)  
  
  # first value  
  total <- sum(head(x, window))  
  res[window] <- total / window  
  
  # remaining values  
  for( i in seq(window+1, n) ) {  
    total <- total + x[i] - x[i-window]  
    res[i] <- total / window  
  }  
  res  
}
```



Hackstucious (hack + astucious) vectorization

```
x <- c(1.3, 3.2, 4.2, 4.5, 6.8)

start <- sum(x[1:3])

head( x, -3 )
1.3 3.2

tail( x, -3 )
4.5 6.8

c( start, start + cumsum( tail(x, -3) - head( x, -3 ) ) )
8.7 11.9 15.5

c( start, start + cumsum( tail(x, -3) - head( x, -3 ) ) ) / 3
2.900000 3.966667 5.166667
```



Comparison

```
library(microbenchmark)
x <- rnorm(1e5)

microbenchmark(
  rollmean1(x, 3),
  rollmean2(x, 3),
  rollmean3(x, 3)
)
Unit: milliseconds
```

	expr	min	lq	mean	median	...
	rollmean1(x, 3)	833.667884	857.507753	971.250098	893.206776	...
	rollmean2(x, 3)	10.539993	11.034244	12.293105	11.396629	...
	rollmean3(x, 3)	1.429817	1.625453	3.070925	3.067068	...

Last observation carried forward





Last observation carried forward

```
na_locf1 <- function(x) {  
  current <- NA  
  
  res <- x  
  for( i in seq_along(x)) {  
    if( is.na(x[i]) ) {  
      # replace with current  
      res[i] <- current  
    } else {  
      # set current  
      current <- x[i]  
    }  
  }  
  
  res  
}
```



Mean carried forward

x	1	3	4	NA	3	2	NA	NA	5
totals	1	4	8	8	11	13	13	13	18
n	1	2	3	3	4	5	5	5	6
				↓			↓	↓	
mean	1	3	4	2.67	3	2	2.6	2.6	5

Mean carried forward

```
na_meancf1 <- function(x) {  
  # ( cumulative sum of non NA values ) / ( cumulative count of non NA )  
  means <- cumsum( replace(x, is.na(x), 0) ) / cumsum(!is.na(x))  
  
  # replace the missing values by the means  
  x[is.na(x)] <- means[is.na(x)]  
  x  
}  
  
# iterative version  
na_meancf2 <- function(x) {  
  total <- 0  
  n <- 0  
  for( i in seq_along(x) ){  
    if( is.na(x[i]) ){  
      x[i] <- total / n  
    } else {  
      total <- x[i] + total  
      n <- n + 1  
    }  
  }  
}
```



Comparisons

```
x <- rnorm(1e5)
x[ sample(1e5, 100) ] <- NA

microbenchmark( na_meancf1(x), na_meancf2(x) )
Unit: milliseconds
```

	expr	min	lq	mean	median	...
na_meancf1(x)		1.176276	2.785667	3.237009	3.474028	...
na_meancf2(x)		16.945271	17.430133	19.678276	18.625274	...



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Auto regressive model

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Auto regressive model, AR

$$X_i = \sum_{j=1}^{np} \phi_j X_{i-j} + \epsilon_i$$

```
ar <- function(n, phi, sd){  
  x <- epsilon <- rnorm(n, sd = sd)  
  np <- length(phi)  
  
  for( i in seq(np+1, n)){  
    x[i] <- sum(x[seq(i-1, i-np)] * phi) + epsilon[i]  
  }  
  x  
}
```

AR in C++

- First `for`, to fill the `np` first values

```
NumericVector x(n) ;

// initial loop
for( ____ ; ____ < np ; ____ ){
    x[i] = R::rnorm(____) ;
}
```

- Main part with outer and inner `for`

```
// outer loop
for( ____ ; ____ ; ____ ){

    double value = rnorm(____) ;

    // inner loop
    for( ____ ; ____ ; ____ ){
        value += ____ ;
    }
    x[i] = value ;
}
```




Moving average simulation

$$X_i = \epsilon_i + \sum_{j=1}^{nq} \theta_j \epsilon_{i-j}$$

```
ma <- function(n, theta, sd){  
  epsilon <- rnorm(n, sd = sd)  
  x <- numeric(n)  
  nq <- length(theta)  
  
  for( i in seq(nq+1, n)){  
    x[i] <- sum(epsilon[seq(i-1, i-nq)] * theta) + epsilon[i]  
  }  
  x  
}
```

Moving average simulation

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

// [[Rcpp::export]]
NumericVector ma( int n, double mu, NumericVector theta, double sd ){
    int nq = theta.size() ;

    // generate the noise vector at once
    // using the Rcpp::rnorm function, similar to the R function
    NumericVector eps = Rcpp::rnorm(n, 0.0, sd) ;

    // init the output vector of size n with all 0.0
    NumericVector x(____) ;

    // start filling the values at index nq + 1
    for( int i=nq+1; i<n; i++){

        _____
    }
    return x ;
}
```



$$\text{ARMA}(p,q) = \text{AR}(p) + \text{MA}(q)$$

$$X_i = \epsilon_i + \sum_{j=1}^{np} \phi_j X_{i-j} + \sum_{j=1}^{nq} \theta_j \epsilon_{i-j}$$



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Congratulations!

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evalCpp and cppFunction

- Evaluating simple C++ statements

```
evalCpp( "40+2" )  
42
```

- Creating a C++ function from the R console

```
cppFunction( "double add( double x, double y){  
    return x + y ;  
}" )  
  
add( 40, 2 )  
42
```



For loops

```
for( init ; condition ; increment ){  
    body  
}
```

- init: what happens at the beginning
- condition: should the loop continue
- increment: after each iteration
- body: what the loop does



For loops

```
for( int i=0; i<n; i++) {  
    // do something with i  
}
```




Vector indexing

```
NumericVector x = ... ;  
int n = x.size() ;
```

```
// first value  
x[0]
```

```
// second value  
x[1]
```

```
// last value  
x[n-1]
```



C++ files with Rcpp

```
#include <Rcpp.h>
```

```
using namespace Rcpp ;
```

```
// [[Rcpp::export]]  
double add( double x, double y) {  
    return x + y ;  
}
```

```
// [[Rcpp::export]]  
double twice( double x) {  
    return 2.0 * x;  
}
```



Typical Rcpp function

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

// [[Rcpp::export]]
double fun( NumericVector x ){

    // extract data from input and prepare outputs
    int n = x.size() ;
    double res = 0.0 ;

    // loop around input and/or output
    for(int i=0; i<n; i++){
        // do something with x[i]
    }

    // return output
    return res ;
}
```



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Congratulations!