Section functions to a smaller domain with section_fun() in the doBy package

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1 Introduction

The **doBy** package contains a variety of utility functions. This working document describes some of these functions. The package originally grew out of a need to calculate groupwise summary statistics (much in the spirit of PROC SUMMARY of the SAS system), but today the package contains many different utilities.

2 Section a functions domain: section_fun()

Let E be a subset of the cartesian product $X \times Y$ where X and Y are some sets. Consider a function f(x,y) defined on E. Then for any $x \in X$, the section of E defined by x (denoted E_x) is the set of y's in Y such that (x,y) is in E, i.e.

$$E_x = \{ y \in Y | (x, y) \in E \}$$

Correspondingly, the section of f(x,y) defined by x is the function f_x defined on E_x given by $f_x(y) = f(x,y)$.

For example, if f(x,y) = x + y then $f_x(y) = f(10,y)$ is a section of f to be a function of y alone.

There are two approaches: 1) replace the section values in the function (default) or 2) store the section values in an auxiliary environment.

2.1 Replace section into function body

Default is to replace section value in functions body:

```
> f <- function(a, b, c=4, d=9){
    a + b + c + d
}
> fr_ <- section_fun(f, list(b=7, d=10))
> fr_

## function (a, c = 4, b = 7, d = 10)
## {
    ## a + b + c + d
## }

> f(a=10, b=7, c=5, d=10)

## [1] 32

> fr_(a=10, c=5)
```

2.2 Using an auxiliary environment

An alternative is to store the section values in an auxiliary environment:

The section values are stored in an extra environment in the scaffold object and the original function is stored in the scaffold functions environment:

```
## $b
## [1] 7
##
## $d
## [1] 10

> ## attr(fe_, "arg_env")$args ## Same result
> get_fun(fe_)

## function(a, b, c=4, d=9){
## a + b + c + d
## }
## <bytecode: 0x556db69d8de0>

> ## environment(fe_)$fun ## Same result
```

3 Example: Benchmarking

Consider a simple task: Creating and inverting Toeplitz matrices for increasing dimensions:

```
> n < -4
> toeplitz(1:n)
         [,1] [,2] [,3] [,4]
                 2
## [1,]
           1
                      2
                            3
## [2,]
           2
                 1
## [3,]
           3
                 2
                      1
                            2
## [4,]
                            1
```

A naive implementation is

```
> inv_toeplitz <- function(n) {
        solve(toeplitz(1:n))
   }
> inv_toeplitz(4)

## [,1] [,2] [,3] [,4]
## [1,] -0.4  0.5  0.0  0.1
```

```
## [2,] 0.5 -1.0 0.5 0.0
## [3,] 0.0 0.5 -1.0 0.5
## [4,] 0.1 0.0 0.5 -0.4
```

We can benchmark timing for different values of n as

```
> library(microbenchmark)
> microbenchmark(
     inv_toeplitz(4), inv_toeplitz(8), inv_toeplitz(16),
     inv_toeplitz(32), inv_toeplitz(64),
     times=5
  )
## Unit: microseconds
##
               expr
                              lq mean median
                      min
                                                  uq
                                                        max neval cld
   inv_toeplitz(4) 114.0 121.3 123.1 124.0 124.2
##
                                                      131.9
                                                                   а
## inv_toeplitz(8) 135.3 140.0 141.3 141.1 144.5 145.5
## inv_toeplitz(16)
                           287.8 314.9 317.0 345.7 355.5
                     268.5
                                                                5
   inv_toeplitz(32)
                    756.0 794.1 1800.9 833.1 953.8 5667.6
                                                                5 ab
## inv_toeplitz(64) 2188.6 2284.3 2748.2 2847.1 3023.1 3398.0
                                                                5
                                                                   b
```

However, it is tedious (and hence error prone) to write these function calls.

A programmatic approach using **section_fun** is as follows: First create a list of sectioned functions:

Each element is a function (a scaffold object, to be precise) and we can evaluate each / all functions as:

```
> fun_list[[1]]

## function (n = 4)

## {

## solve(toeplitz(1:n))

## }

> fun_list[[1]]()

## [,1] [,2] [,3] [,4]
```

```
## [1,] -0.4 0.5 0.0 0.1

## [2,] 0.5 -1.0 0.5 0.0

## [3,] 0.0 0.5 -1.0 0.5

## [4,] 0.1 0.0 0.5 -0.4
```

To use the list of functions in connection with microbenchmark we bequote all functions using

```
> bquote_list <- function(fnlist){
    lapply(fnlist, function(g) {
        bquote(.(g)())
    }
    )
}</pre>
```

We get:

```
> bq_fun_list <- bquote_list(fun_list)</pre>
> bq_fun_list[[1]]
## (function (n = 4)
## {
##
       solve(toeplitz(1:n))
## })()
> ## Evaluate one:
> eval(bq_fun_list[[1]])
        [,1] [,2] [,3] [,4]
## [1,] -0.4 0.5 0.0 0.1
## [2,] 0.5 -1.0 0.5 0.0
## [3,] 0.0 0.5 -1.0 0.5
## [4,] 0.1 0.0 0.5 -0.4
> ## Evaluate all:
> ## sapply(bq_fun_list, eval)
```

To use microbenchmark we must name the elements of the list:

```
> names(bq_fun_list) <- n.vec
> microbenchmark(
    list = bq_fun_list,
    times = 5
)
```

```
uq
##
   expr min lq mean median
                                        max neval cld
##
        114.4
              116.4
                     122.2 124.4
                                 126.2
                                        129.9
                                                  5 a
##
        131.2
               144.0
                     148.0
                           145.2
                                  153.5
                                        166.0
        237.0 252.2
                                  331.2 679.4
                     364.3 321.7
                                                  5 a
##
     32 709.8 964.9 990.1 973.2 989.0 1313.5
                                                  5 b
##
     64 2144.8 2226.1 2500.1 2409.8 2518.8 3201.1
                                                  5
```

To summarize: to experiment with many difference values of n we can do