

# *Debug and profiling in R*

## ANF R

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## Roadmap

### Simple rules

Herb Sutter

Agile manifesto & co

### Debug

Principle

Anticipating bugs

Tracking bugs with appropriate tools - pure R code

Tracking bugs with appropriate tools - R/C++ code

### Profiling

Principle

Recording time and space requirements

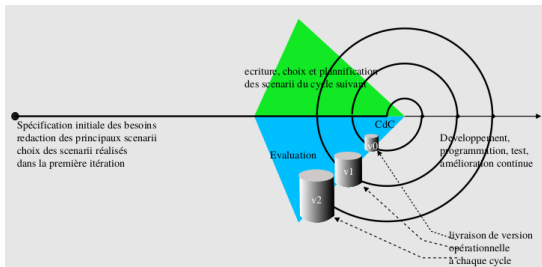
Performance measurements with appropriate tools - pure R code

Performance measurements with appropriate tools - R/C++ code

- ▶ Correctness (**this course**), simplicity and clarity come first
- ▶ Don't optimize prematurely (**this course**)
- ▶ Give one entity one cohesive responsibility
- ▶ Use a version control system

## └ Simple rules

## └ Agile manifesto &amp; co



- ▶ Repeat very short development cycles
- ▶ Write tests first (this course, quickly)
- ▶ Adopt naming conventions and use explicit names
- ▶ Don't hesitate to refactor your code (this course)
- ▶ Submit frequent deliveries

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*Finding a bug is a process of confirming the many things that you believe are true - until you find one which is not true. (Norm Matloff)*

1. Realise you have a bug
2. Make it repeatable
3. Figure out where it is
4. Fix it and test



1. Tests
2. Tests
3. Debugger
4. Tests

Print-like method : `stop()`, `message()`, `warning()`

Condition handling method : `try()`, `tryCatch()`

Defensive programming or "failing fast", i.e. never suppose anything

## Debug

### Tracking bugs with appropriate tools - pure R code

- ▶ Determining the sequence of calls that lead to an error where
- ▶ Consulting the content of variables print

Global Environment	
<b>Data</b>	
distance.matrix	Large matrix (1000000 elements, 7.7 Mb)
points	num [1:1000, 1:2] 0.3748 0.9824 0.425 0.677 0.0518 ...
<b>Values</b>	
clusters	List of 4
N	1000
x	num [1:250] 0.3748 0.9824 0.425 0.677 0.0518 ...
y	num [1:250] 0.609 0.623 0.789 0.701 0.301 ...

- ▶ Setting breakpoints break
- ▶ Execute the code step-by-step next, step



TP Exercise(s) 1



└ Debug

└ Tracking bugs with appropriate tools - R/C++ code

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Much more complicated...

▶ [Bioconductor guidelines](#)

DEMO gdb

DEMO valgrind

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*Programmers waste enormous amounts of time thinking about, or worrying about, the speed of noncritical parts of their programs. (Donald Knuth)*

*Writing faster but incorrect code... Writing code that you think faster, but is actually no better. (Hadley Wickam)*

1. Find the bottleneck
2. Refactor the code
3. Consider using third party libraries or packages
4. Loop to 1



1. Profiler
2. Advanced R or Rcpp
3. Web browser, support from R-help, calcul-list or colleagues, forums
4. Loop to 1

## └ Profiling

└ Recording time and space requirements

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Print-like method : `sys.time()`, `microbenchmark()`, `mem_used()`, `top` (Unix only)

It is mandatory to have a methodology for measuring performance : small to large datasets or problem dimensions + estimation of the theoretical complexity

TP Exercise(s) 2,3

- Profiling

- Performance measurements with appropriate tools - pure R code

R proposes a *sampling* profiler called *Rprof*: it stops the execution of the code every few milliseconds and writes in a file which function is currently executing.

For *Rprof* in parallel, the workers (not the master) launch *Rprof* and specify different files.

Now, let's understand the *call graph*!

```
by.self
      self.time self.pct total.time total.pct
".C"          35.60  32.23    35.60   32.23
"data.frame"  10.36   9.38    59.32   53.70
"deparse"     6.96   6.30    23.74   21.49
"FUN"         6.38   5.78   110.46  100.00
".deparseOpts" 6.20   5.61    9.70    8.78
"as.data.frame.numeric" 4.94   4.47   32.08   29.04
"as.data.frame" 3.94   3.57   36.02   32.61
"nmatch"      3.22   2.92   14.94   13.53
"match"       3.16   2.86    3.50    3.17
"as.list"     2.92   2.64    3.80    3.49
"length"      2.54   2.30    2.54    2.30
"make.names"  2.50   2.26    2.80    2.59
"mode"        2.10   1.90   13.06   11.82
"paste"       2.06   1.86   25.80   23.36
"row.names"   1.80   1.63    2.54    2.30
"unlist"      1.58   1.43    3.12    2.82
"meanRuptr_c" 1.34   1.21   22.88   20.71
".row_names_info" 1.30   1.18    1.30    1.18
".set_row_names" 1.20   1.09    1.20    1.09
"names"       1.10   1.00    1.10    1.00
"%>%>"        1.06   0.96   15.46   14.00
```

```
by.total
      total.time total.pct self.time self.pct
"FUN"          110.46  100.00    6.38    5.78
"lapply"       110.46  100.00    0.00    0.00
"multiseg"     110.46  100.00    0.00    0.00
"multisegmean" 110.46  100.00    0.00    0.00
"standardGeneric" 110.46  100.00    0.00    0.00
"sapply"       95.54   86.49    0.00    0.00
"multisegout"  92.78   83.99    0.02    0.02
"data.frame"   59.32   53.70   10.36   9.38
"as.data.frame" 36.02   32.61    3.94   3.57
".C"           35.60   32.23   35.60   32.23
"as.data.frame.numeric" 32.08   29.04    4.94   4.47
"force"        26.16   23.68    0.36   0.33
"paste"        25.80   23.36    2.06   1.86
"deparse"      23.74   21.49    6.96   6.30
"meanRuptr_c" 22.88   20.71    1.34   1.21
"%>%>"        15.46   14.00    1.06   0.96
"unisegmean"   15.22   13.78    0.00   0.00
"segmeanCO"    15.10   13.67    0.00   0.00
"match"        14.94   13.53    3.22   2.92
"collbrLR_c"   14.72   13.33    0.00   0.00
"mode"         13.06   11.82    2.10   1.90
```

TP Exercise(s) 4,5,6

See forums or read "Writing R extensions" (that suggests the use of oprofile)

```
samples % symbol name
81898 19.0842 Liste::resetMaillonBorders(Polynome2*)
49520 15.2678 Polynome2::roots(double)
38476 11.8628 Polynome2::minOrMax(double*, double*, int*)
33504 10.3298 Liste::removeDoublon()
17497 5.3946 Polynome2::add(double, double, double)
16009 4.9358 Liste::getPolynome()
9972 3.0745 Liste::computeMinOrMax(double*, int*)
9279 2.8609 Liste::add(double, double, double)
9141 2.8183 Liste::getNext()
7334 2.2612 colibri_c(double*, int*, int*, double*, double*, int*, double*)
6953 2.1437 Liste::resetAllBorders(Polynome2*)
6241 1.9242 Polynome2::reset(double, double, double, int)
5754 1.7741 Liste::computeRoots(double)
5413 1.6689 Polynome2::delta(double)
5398 1.6643 Liste::checkForDoublon()
5386 1.6606 Polynome2::getRacine2()
4740 1.4614 Liste::insert(Liste*)
4116 1.2690 Polynome2::setStatus(int)
```

DEMO oprofile