Scalability

Preparing for Production

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Scalability and Production

When put into production code gets used more and on more data.

We will likely have to consider scalability of our methods in

- Computation time
- Memory requirements

When doing so we have to balance a trade-off between development costs and usage costs.

Example: Bayesian Inference

- MCMC originally takes ~24 hours
- Identifying and amending bottlenecks in code reduced this to ~24 minutes.

Is this actually better?

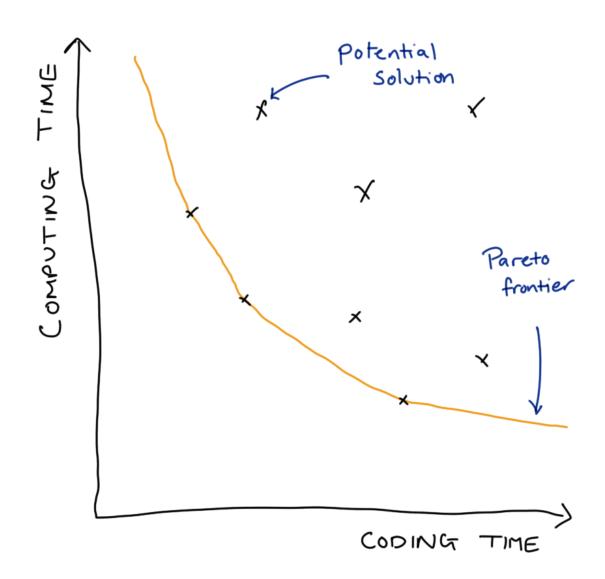
- human hours invested
- frequency of use

- safe / stable / general / readable
- trade for scalability

Knowing when to worry

Sub-optimal optimisation can be worse than doing nothing

... programmers have spent far too much time worrying about efficiency in the wrong places and at the wrong times; premature optimisation is the root of all evil (or at least most of it) in programming. - Donald Knuth



This Lecture

- Basic profiling to find bottlenecks.
- Some simple solutions
- Strategies for scalable (R) code
- Signpost advanced methods & further reading

Profiling your code: basics

R as a stopwatch

```
1 t_start <- Sys.time()
2 Sys.sleep(0.5) # YOUR CODE
3 t_end <- Sys.time()
4
5 t_end - t_start</pre>
```

Time difference of 0.5098979 secs

```
1 library(tictoc)
2
3 tic()
4 Sys.sleep(0.5) # YOUR CODE
5 toc()
```

0.505 sec elapsed

With {tictoc} we can get fancy

```
1 tic("total")
2 tic("first, easy part")
3 Sys.sleep(0.5)
4 toc(log = TRUE)
5 ## first, easy part: 0.505 sec elapsed
6 tic("second, hard part")
7 Sys.sleep(3)
8 toc(log = TRUE)
9 ## second, hard part: 3.008 sec elapsed
10 toc()
11 ## total: 3.519 sec elapsed
```

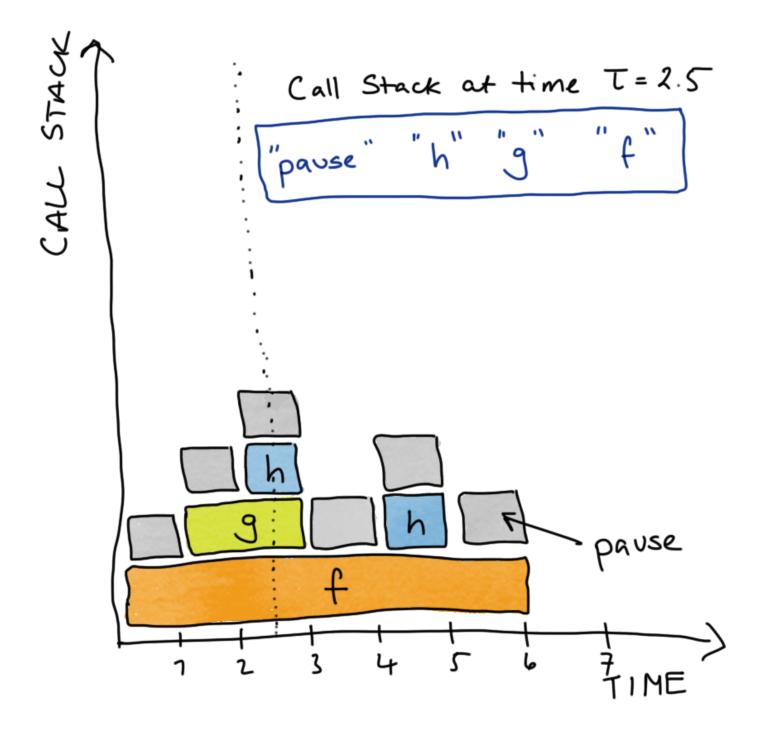
Profiling your code in detail

To diagnose scaling issues you have to understand what your code is doing.

- Stop the code at time τ and examine the call-stack.
 - The current function being evaluated, the function that called that, the function that called that, ..., top level function.
- Do this a lot and you can measure (estimate) the proportion of working memory (RAM) uses over time and the time spent evaluating each function.

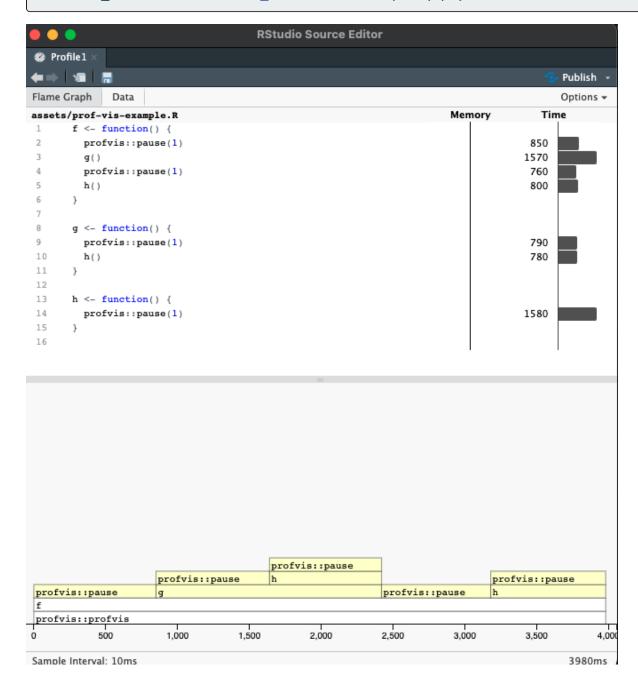
Profiling: Toy Example

```
h <- function() {</pre>
      profvis::pause(1)
 3 }
   g <- function() {</pre>
      profvis::pause(1)
      h()
8 }
 9
10 f <- function() {</pre>
      profvis::pause(1)
12
      g()
13
      profvis::pause(1)
14
      h()
15 }
```



Profiling: How To

- 1 source("assets/prof-vis-example.R")
- 2 profvis::profvis(f())



Notes on Time Profiling

- Will get slightly different results each time you run the function
 - Changes to internal state of computer
 - Usually not a big deal, mainly effects fastest parts of code
 - Be careful with stochastic simulations
 - Use set.seed() to make a fair comparison over many runs.

Notes on Profiling

Function Source

```
1 pad_with_NAs
function(x, n_left, n_right){
  c(rep(NA, n_left), x, rep(NA, n_right))
}
```

Compiled functions have no R source code.

Compiled Function

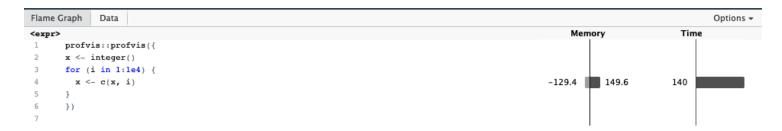
```
1 mean
function (x, ...)
UseMethod("mean")
<bytecode: 0x7fabbac6d718>
<environment: namespace:base>
```

 Profiler does not extend into compiled code, see {jointprof} if you really need this.

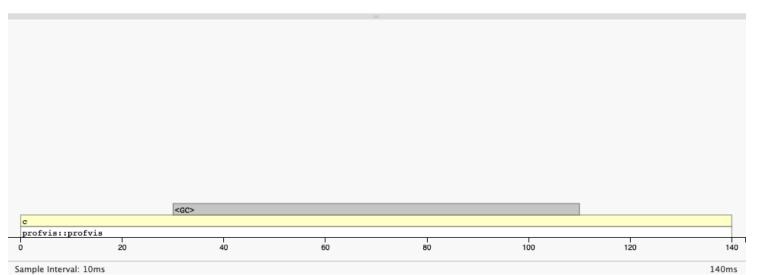
Memory Profiling

profvis () can similarly measure the memory usage of your code.

```
1 x <- integer()
2 for (i in 1:1e4) {
3    x <- c(x, i)
4 }</pre>
```



- Copy-on-modify behaviour makes growing objects slow.
- Pre-allocate storage where possible.
- Strategies and structures, see R inferno and Effecient R.



Tips to work at scale

Vectorise

```
1 x <- 1:10
2 y <- 11:20
3 z <- rep(NA, length(x))
4
5 for (i in seq_along(x)) {
6  z[i] <- x[i] * y[i]
7 }</pre>
```

```
1 x <- 1:10
2 y <- 11:20
3 z <- x * y
```

Use and write functions with vectorised inputs.

```
1 \text{ rnorm(n = 100, mean = 1:10, sd = rep(1, 10))}
```

Be careful of recycling!

Special vectors: Linear Algebra

```
1 X <- diag(x = c(2, 0.5))
2 y <- matrix(data = c(1, 1), ncol = 1)
3
4 X %*% y

[,1]
[1,] 2.0
[2,] 0.5</pre>
```

More on vectorising: Noam Ross Blog Post

For loops in disguise: the apply family

Functional programming equivalent of a for loop. [apply(), mapply(), lapply(),...]

Apply a function to each element of a list-like object.

```
1 # MARGIN = 1 => rows, MARGIN = 2 => columns
2 apply(X = A, MARGIN = 1, FUN = sum)
[1] 22 26 30
```

Generalises functions from {matrixStats}

```
1 rowSums(A)
[1] 22 26 30
```

For loops in disguise: purrr::map

Iterate over a single object with map ():

```
1 mu <- c(-10, 0, 10)
2 purrr::map(.x = mu, .f = rnorm, n = 5)

[[1]]
[1] -10.630941 -10.536954 -10.393322 -8.395591 -8.770291

[[2]]
[1] -0.2173831  0.4013440 -2.2716535 -1.5199591  0.2817534

[[3]]
[1] 8.217122  7.209396  9.051645  10.628293  11.263494</pre>
```

Iterate over multiple objects map2() and pmap():

```
1  mu <- c(-10, 0, 10)
2  sigma <- c(0, 0.1, 0)
3  purrr::map2(.x = mu, .y = sigma, .f = rnorm, n = 5)

[[1]]
[1] -10 -10 -10 -10 -10

[[2]]
[1] -0.097381610 -0.104544473 -0.063586494 -0.002777205
-0.105194011

[[3]]
[1] 10 10 10 10 10 10</pre>
```

For more details and variants see Advanced R chapters 9-11 on functional programming.

Easy parallelisation with furrr

- {parallel} and {futures} allow parallel coding over multiple cores.
- Powerful, but steep learning curve.
- {furrr} makes this very easy, just add future_ to purrr verbs.

Need to be very careful handling RNG. See R-bloggers for more details.

Sometimes R doesn't cut it



- An API for running C++ code in R
 - Loops that need to be run in order
 - Lots of function calls (e.g. deep recursion)
 - Fast data structures
- Beyond our scope but good to know exists. Starting point: Advanced R Chapter 25.

Wrapping up Summary

- 1. Pick you battles wisely
- 2. Target your energy with profiling
- 3. Scale loops with vectors
- 4. Scale loops in parallel processing
- 5. Scale in another language

Help!

- Articles and blog links
- The R inferno (Circles 2-4)
- Advanced R (Chapters 23-25),
- Efficient R (Chapter 7).

