# Parallel computations with $\ensuremath{\mathbb{R}}$

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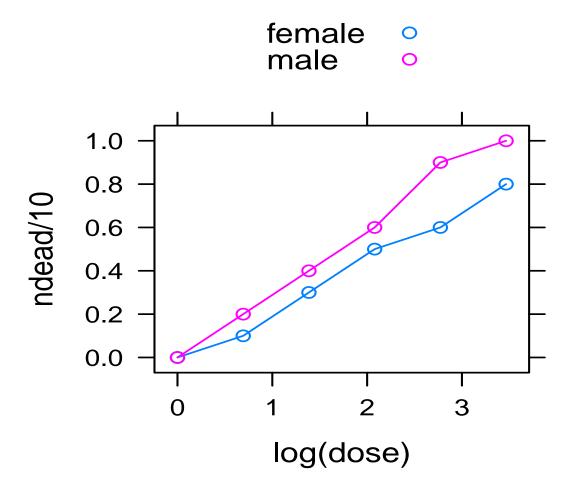
# 1 Introduction

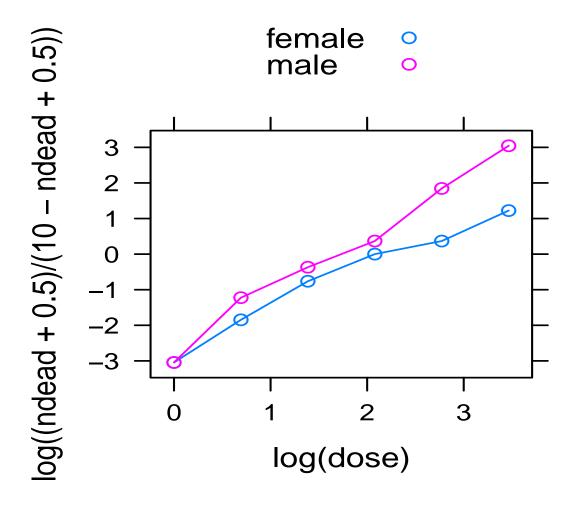
- For some years now, computers have not become much faster (it seems that in practice there is an upper limit of about 3 GHz).
- Instead, most (all?) computers today come with multiple cores and hence the ability to make parallel computations.
- There are various R packages for parallel computations. The **parallel** package is shipped with the R distribution and this is the package we illustrate here.

# 2 Parametric bootstrap

# 2.1 A logistic regression

```
R> budworm
      sex dose ndead ntotal
     male
                    0
                          10
     male
                          10
     male
                          10
     male
                    6
                          10
     male
            16
                    9
                          10
     male
            32
                   10
                          10
   female
                    0
                          10
  female
                          10
   female
             4
                    3
                          10
10 female
             8
                    5
                          10
11 female
            16
                    6
                          10
12 female
            32
                    8
                          10
```





Very vanilla logistic regression. Is there an effect of sex?

### 2.2 Parametric bootstrap

Alternative to  $-2 \log Q$  test: Parametric bootstrap – seems attractive when large sample asymptotics is questionable:

- Compare two models  $M_1$  and  $M_0$  where  $M_0 \subset M_1$ .
- Fit both models to data; Gives  $\hat{\theta}_1$ ,  $\hat{\theta}_0$  and  $t_{obs}$  (test statistic)
- Draw B parametric bootstrap sample datasets  $y^1, \ldots y^B$  from  $p_0(\cdot|\hat{\theta}_0)$ . For each simulated dataset  $y^b$ , calculate test statistic  $t^b$ .
- Evaluate how extreme  $t_{obs}$  is in  $\{t^1, \dots t^B\}$ .

### 2.3 Non-parallel version

Need to refit glm to new response variable:

Need to do so many times and calculate reference distribution:

```
R> pboot <- function(lg, sm, nsim=10){
+    simdata <- simulate(sm, nsim)
+    ref <- rep.int(NA,nsim)
+    for (ii in 1:nsim){
+        y.new <- simdata[,ii]
+        ref[ii] <- 2*(logLik(refit_glm(lg, y.new))-logLik(refit_glm(sm, y.new)))
+    }
+    ref
+  }</pre>
```

```
R> anova(mm1, mm0, test="Chisq")
Analysis of Deviance Table
Model 1: cbind(ndead, ntotal - ndead) ~ sex + logdose
Model 2: cbind(ndead, ntotal - ndead) ~ logdose
  Resid. Df Resid. Dev Df Deviance Pr(>Chi)
        9
                5.0466
1
        10 8.7883 -1 -3.7417 0.05307 .
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
R> set.seed(123)
R> Nsim <- 2000
R > (obsdev \leftarrow c(2*(logLik(mm1)-logLik(mm0))))
[1] 3.741666
R> system.time({    refdev <- pboot(mm1, mm0, nsim=Nsim) })</pre>
   user system elapsed
  10.51
           0.02 10.55
R> sum(refdev > obsdev) / length(refdev)
[1] 0.0505
```

### 2.4 Parallel version

To do parallel computations, first detect the number of cores available. Next create a handle on the clusters:

```
R> library("parallel")
R> ## Number of cores:
R> (nc <- detectCores())
[1] 8
R> ## Create clusters
R> cl <- makeCluster(rep("localhost", nc))</pre>
```

When done with the parallel computations, close the clusters before leaving R:

```
R> ## Remember to shut down clusters before quitting R
R> stopCluster(cl)
```

```
R> t0 <- proc.time()
R> Nsim <- 2000
R> (nsim2 <- round(Nsim / nc))

[1] 250

R> ## Export global environment to each cluster
R> clusterExport(cl, ls(envir=.GlobalEnv), envir = .GlobalEnv)
R> ## Spec for random number generator
R> clusterSetRNGStream(cl)
R> ## Time to get things up and running
R> proc.time()-t0

user system elapsed
0.02 0.03 0.05
```

#### Notice:

- We get a substantial saving in computing time, but not by a factor equal to the number of cores.
- There is some overhead in setting things up for parallel computations.

# 3 Matrix multiplication

Multiplication of a  $p \times q$  and a  $q \times r$  matrix has complexity of the order pqr.

```
R> nr <- 3
R> (A <- matrix(round(rnorm(nr^2),1),nr=nr))</pre>
     [,1] [,2] [,3]
[1,] -0.2 0.4 -1.6
[2,] 0.8 0.2 0.5
[3,] -1.2 1.0 0.3
R > (B < -t(A) + 4)
     [,1] [,2] [,3]
[1,] 3.8 4.8 2.8
[2,] 4.4 4.2 5.0
[3,] 2.4 4.5 4.3
R> A %*% B
      [,1] [,2] [,3]
[1,] -2.84 -6.48 -5.44
[2,] 5.12 6.93 5.39
[3,] 0.56 -0.21 2.93
```

Idea: Split A by rows, do multiplications and stack the results:

```
R> A[1,,drop=FALSE] %*% B
      [,1] [,2] [,3]
[1,] -2.84 -6.48 -5.44
R> A[2,,drop=FALSE] %*% B
     [,1] [,2] [,3]
[1,] 5.12 6.93 5.39
R> A[3,,drop=FALSE] %*% B
     [,1] [,2] [,3]
[1,] 0.56 -0.21 2.93
R> rbind(
+ A[1,,drop=FALSE] %*% B,
+ A[2,,drop=FALSE] %*% B,
+ A[3,,drop=FALSE] %*% B)
      [,1] [,2] [,3]
[1,] -2.84 -6.48 -5.44
[2,] 5.12 6.93 5.39
[3,] 0.56 -0.21 2.93
```

```
R> ## Use 2 clusters
R> c12 <- c1[1:2]
R> (idx <- splitIndices(nrow(A), length(c12)))</pre>
[[1]]
[1] 1
[[2]]
[1] 2 3
R> (Alist <- lapply(idx, function(ii) A[ii,,drop=FALSE]))</pre>
[[1]]
    [,1] [,2] [,3]
[1,] -0.2 0.4 -1.6
[[2]]
     [,1] [,2] [,3]
[1,] 0.8 0.2 0.5
[2,] -1.2 1.0 0.3
```

```
R> ans <- clusterApply(c12, Alist, function(aa, BB) aa %*% BB, B)
R> do.call(rbind, ans)

[,1] [,2] [,3]
[1,] -2.84 -6.48 -5.44
[2,] 5.12 6.93 5.39
[3,] 0.56 -0.21 2.93

R> A %*% B

[,1] [,2] [,3]
[1,] -2.84 -6.48 -5.44
[2,] 5.12 6.93 5.39
[3,] 0.56 -0.21 2.93
```

### 3.1 Parallel version

Define parallel matrix multiplication function:

#### Notice:

```
R> get("%*%")
function (x, y) .Primitive("%*%")
```

```
R > nr < -5
R> A <- matrix(round(rnorm(nr^2),1),nr=nr)</pre>
R > B < -t(A) + 4
R> matprod.par(cl, A, B)
      [,1] [,2] [,3] [,4] [,5]
[1,] -1.65 -2.09 -6.30 -3.94 -5.51
[2,] -0.49 6.05 -1.91 -3.04 -1.22
[3,] 4.90 7.69 12.41 5.21 10.40
[4,] 13.26 12.56 11.21 17.60 12.50
[5,] 2.09 4.78 6.80 2.90 6.30
R> A %*% B
      [,1] [,2] [,3] [,4] [,5]
[1,] -1.65 -2.09 -6.30 -3.94 -5.51
[2,] -0.49 6.05 -1.91 -3.04 -1.22
[3,] 4.90 7.69 12.41 5.21 10.40
[4,] 13.26 12.56 11.21 17.60 12.50
[5,] 2.09 4.78 6.80 2.90 6.30
```

# 4 The heat equation

For a function u(x,y,t) of spatial variables (x,y) and time variable t, the heat equation is

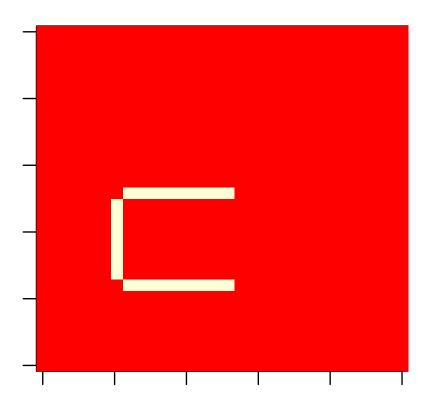
$$\frac{\partial u}{\partial t} - \alpha \left( \frac{\partial^2 u}{\partial x^2} + \frac{\partial^2 u}{\partial y^2} \right) = 0, \quad \alpha > 0$$

This equation describes how the temperature u(x, y, t) at location (x, y) changes over time t as heat spreads in space.

To solve this equation we must speficify the initial conditions u(x,y,0) and a set of boundary conditions.

Suppose we throw a heated horse shoe (with temperature 1) on a cold plate (with temperature 0):

```
R> nr <- nc <- 30
R> mm <- matrix(0,nrow=nr, ncol=nc)
R> mm[7,9:15] <- mm[8:16,16] <- mm[8:16,8] <- 1
R> image(mm)
```



The temperature at the boundaries (outermost rows and columns) is assumed to remain constantly equal to zero. (Called a Dirichlet condition).

On a regular grid, this equation can be solved numerically as

$$u(x, y, t + 1) = u(x, y, t) +$$

$$\alpha(u(x + 1, y, t) + u(x - 1, y, t) - 2u(x, y, t)) +$$

$$\alpha(u(x, y + 1, t) + u(x, y - 1, t) - 2u(x, y, t))$$

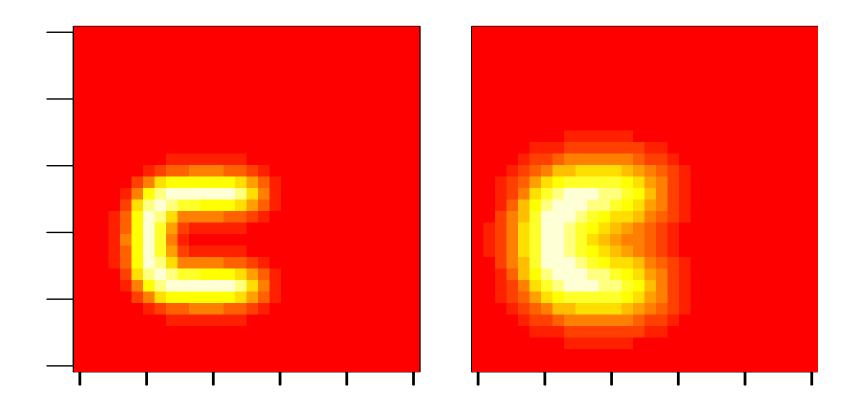
```
R> ## Update in one time step
R> heat <- function(mm, cx=.1, cy=.1, nr=nrow(mm), nc=ncol(mm)){
     mm2 < - mm
     for (ii in 2:(nr-1)){
       for (jj in 2:(nc-1)){
         mm2[ii,jj] <- mm[ii,jj] +
           cy * (mm[ii-1,jj]+mm[ii+1,jj]-2*mm[ii,jj]) +
             cx * (mm[ii,jj-1]+mm[ii,jj+1]-2*mm[ii,jj])
     7
     mm2
R> library(compiler) ## Byte compiling gives a factor 5 in speed
R> heat.cmp <- cmpfun(heat)</pre>
R> ## Iterate over time
R> heat.iter <- function(mm, n=1, cx=.1, cy=.1){
     for (ii in 1:n){
       mm <- heat.cmp(mm, cx, cy)</pre>
     }
     mm
```

```
R> mm.10 \leftarrow heat.iter(mm, n=10)
```

R> mm.30 <- heat.iter(mm.10,n=20)

R> image(mm.10)

R> image(mm.30,yaxt='n')



### 4.1 Numeric estimation of second derivatives

Suppose that we want to estimate the second derivative of a function f at a point x and that we know the values of the function at three points

$$(x-h, f(x-h)), (x, f(x)), (x+h, f(x+h))$$

Since f''(x) is the derivative of the function f' at the point x we can estimate f''(x) by

$$f''(x) \approx \frac{f'(x+\frac{h}{2}) - f'(x-\frac{h}{2})}{h}$$

We can estimate f'(x-h/2) and f'(x+h/2) by

$$f'(x - \frac{h}{2}) \approx \frac{f(x) - f(x - h)}{h}$$
 $f'(x + \frac{h}{2}) \approx \frac{f(x + h) - f(x)}{h}$ 

Putting this together gives

$$f''(x) \approx \frac{\frac{f(x+h)-f(x)}{h} - \frac{f(x)-f(x-h)}{h}}{h}$$

$$= \frac{f(x+h)-2f(x)+f(x-h)}{h^2}$$

Partial derivatives can be estimated the same way:

$$\frac{\partial^2}{\partial x^2} f(x,y) \approx \frac{f(x+h,y) - 2f(x,y) + f(x-h,y)}{h^2}$$

$$\frac{\partial^2}{\partial y^2} f(x,y) \approx \frac{f(x,y+h) - 2f(x,y) + f(x,y-h)}{h^2}$$

# 5 EXERCISES: The heat equation

The update over time for the heat equation is an obvious candidate for parallel computing:

- 1. Divide the grid into regions for example by splitting by the rows.
- 2. Update the regions.
- 3. Put the updated regions together and make the necessary modifications on the boundary between the regions.

#### Exercise:

1. Implement a parallel version of the algorithm for updating the heat equation.