

Calcul parallèle avec R

ANF R

Vincent Miele
CNRS

07/10/2015

Pour chaque exercice, il est nécessaire d'ouvrir une fenêtre de visualisation des processus (`Terminal + top` sous Linux et Mac OS X, `Gestionnaire des tâches` sous Windows) et d'observer le comportement des processus R.

Par ailleurs, si votre machine a plus que 4 GB de RAM, remplacer dans ce qui suit `m=3000` par `m=5000`. Enfin, avant chaque exercice, il est préférable de relancer R (sans sauvegarde de l'environnement).

Exercice 1 (Parallel apply)

1. tester la version séquentielle

```
n=1000
m=30000
M = matrix(rnorm(n*m), ncol=m)
V = matrix(rnorm(10*m), ncol=m)

eucd <- function(u){
  euc.distu = apply(V, 1, function(w) sqrt(sum((u - w)^2)))
}

gc()
ptm <- proc.time()
euc.dist1 = apply(M, 1, eucd)
proc.time() - ptm
```

2. tester la version *snow-like*

```
library(parallel)

gc()
cl <- makeCluster(2, type="PSOCK")
clusterExport(cl, list("V"))
ptm <- proc.time()
euc.dist2 = parApply(cl, M, 1, eucd)
proc.time() - ptm
stopCluster(cl)
```

3. tester la version *multicore-like*

```
gc()
```

```

cl <- makeCluster(2, type="FORK")
ptm <- proc.time()
euc.dist3 = parApply(cl, M, 1, eucd)
proc.time() - ptm
stopCluster(cl)

```

Exercice 2 (Parallel lapply)

1. tester la version séquentielle

```

n=1000
m=30000
L = lapply(1:n, rnorm, n = m )
V = matrix(rnorm(10*m), ncol=m)

eucd <- function(u){
  euc.distu = apply(V, 1, function(w) sqrt(sum((u - w)^2)))
}

gc()
ptm <- proc.time()
euc.dist1 = lapply(L, eucd)
proc.time() - ptm

```

2. tester la version *snow-like*

```

library(parallel)
gc()
cl <- makeCluster(2, type="PSOCK")
clusterExport(cl, list("V"))
ptm <- proc.time()
euc.dist2 = parLapply(cl, L, eucd)
proc.time() - ptm
stopCluster(cl)

```

3. tester les deux versions *multicore-like*

```

gc()
ptm <- proc.time()
euc.dist1 = mclapply(L, eucd, mc.cores=2)
proc.time() - ptm

gc()
cl <- makeCluster(2, type="FORK")
ptm <- proc.time()
euc.dist3 = parLapply(cl, L, eucd)
proc.time() - ptm
stopCluster(cl)

```

Exercice 3 (mcparallel et mccollect)

Tester le lancement de tâches (Linux et Mac OS X seulement)

```
1. library(parallel)
   library(igraph)

   mis <- function(g){
     ptm <- proc.time()
     lis = largest.independent.vertex.sets(g)[1]
     t = proc.time() - ptm
     cat("Graph with ",vcount(g)," vertices : DONE in ",as.numeric(t[3]),"\n")
     lis
   }

   g1 <- erdos.renyi.game(70, 1/25)
   mis1 = mcparallel(mis(g1))
   mccollect(mis1, wait=FALSE)

   g2 <- erdos.renyi.game(60, 1/25)
   mis2 = mcparallel(mis(g2))
   mccollect(list(mis1, mis2), wait=TRUE)
```

Exercice 4 (Parallel foreach)

1. tester la version séquentielle

```
library(foreach)
n=1000
m=30000
M = matrix(rnorm(n*m), ncol=m)
V = matrix(rnorm(10*m), ncol=m)

ptm <- proc.time()
euc.dist4 = foreach(i=1:n, .combine=c) %do% eucd(M[i,])
proc.time() - ptm
```

2. tester tester la version *snow-like*

```
library(foreach)
library(parallel)
library(doParallel)
cl <- makeCluster(2)
registerDoParallel(cl)
ptm <- proc.time()
euc.dist6 = foreach(i=1:n, .combine=c) %dopar% eucd(M[i,])
proc.time() - ptm
stopCluster(cl)
```

3. tester tester la version *multicore-like*

```
registerDoParallel(cores=2)
ptm <- proc.time()
euc.dist5 = foreach(i=1:n, .combine=c) %dopar% eucd(M[i,])
proc.time() - ptm
```

Exercice 5 (Scheduling)

1. un peu d'initialisation...

```
library(parallel)

lazy <- function(i){
  Sys.sleep(i)
  cat(Sys.getpid()," waited ",i," seconds\n")
  Sys.getpid()
}

L = list(5,30,5,10,5,10)
```

2. comparer les 2 versions *multicore-like* (Linux et Mac OS X)

```
ptm <- proc.time()
res = mclapply(L, lazy, mc.cores=2) # preschedule 1 by 1
proc.time() - ptm
res

ptm <- proc.time()
res = mclapply(L, lazy, mc.cores=2, mc.preschedule = FALSE)
proc.time() - ptm
res
```

3. comparer les 2 versions *snow-like*

```
cl <- makeCluster(2)
clusterSplit(cl, L)
ptm <- proc.time()
res = parLapply(cl, L, lazy) # preschedule by chunks
proc.time() - ptm
stopCluster(cl)
res

cl <- makeCluster(2)
ptm <- proc.time()
res = parLapplyLB(cl, L, lazy) # fake, watch the source
proc.time() - ptm
stopCluster(cl)
res
```

4. reprendre avec une "vrai" function parLapplyLB définie sur <http://examples.oreilly.com/0636920021421/snow/loadbalancing.R>

```
parLapplyLB <- function(cl, x, fun, ...) {
  clusterCall(cl, LB.init, fun, ...)
  r <- clusterApplyLB(cl, x, LB.worker)
  clusterEvalQ(cl, rm('.LB.fun', '.LB.args', pos=globalenv()))
  r
}
LB.init <- function(fun, ...) {
  assign('.LB.fun', fun, pos=globalenv())
  assign('.LB.args', list(...), pos=globalenv())
```

```

NULL
}
LB.worker <- function(x) {
do.call('.LB.fun', c(list(x), .LB.args))
}

```

Exercice 6 (Scheduling for real)

1. un peu d'initialisation... (notamment tirage aléatoire de graphes)

```

library(parallel)
library(igraph)

L = lapply(list(50,10,50,25,50,25), function(i){g <- erdos.renyi.game(70, 1/i)})

mis <- function(g){
ptm <- proc.time()
lis = largest.independent.vertex.sets(g)[1]
t = proc.time() - ptm
cat("Graph with ",vcount(g)," vertices : DONE in ",as.numeric(t[3]),"\n")
lis
}

```

2. comparer les 2 versions *multicore-like* (Linux et Mac OS X)

```

ptm <- proc.time()
res = mclapply(L, mis, mc.cores=2)
proc.time() - ptm

ptm <- proc.time()
res = mclapply(L, mis, mc.cores=2, mc.preschedule = FALSE)
proc.time() - ptm

```

3. comparer les 2 versions *snow-like*

```

cl <- makeCluster(2)
ptm <- proc.time()
clusterEvalQ(cl, {library(igraph); NULL})
res = parLapply(cl, L, mis) # preschedule by chunks
proc.time() - ptm
stopCluster(cl)

cl <- makeCluster(2)
ptm <- proc.time()
clusterEvalQ(cl, {library(igraph); NULL})
res = parLapplyLB(cl, L, mis) # fake, watch the source
proc.time() - ptm
stopCluster(cl)

```

4. éventuellement reprendre depuis le début de l'exercice pour tirer de nouveaux graphes

Exercice 7 (clusterExport)

Tester la présence et l'absence du clusterExport

```
library(parallel)

n=1000
m=30000
M <- matrix(rnorm(n*m), ncol=m) # M is in GlobalEnv
cl <- makeCluster(2, type="PSOCK")
doSleep1 <- function(i) {Sys.sleep(i); M[1,1]=0;}
# clusterExport(cl, list("M"))
res = parLapply(cl, list(20,20), doSleep1)
stopCluster(cl)
```

Exercice 8 (Serialization et mal de tête)

1. après avoir relancé R, observer la consommation mémoire des processus au fur et à mesure de l'exécution de la fonction parSleep. Comprendre le contenu des fichiers de sorties output_xxx.txt.

```
library(parallel)

parSleep <- function(){
  n=1000
  m=30000
  M <- matrix(rnorm(n*m), ncol=m) # M is not in GlobalEnv
  # then clusterExport is impossible
  M2 <- matrix(rnorm(n*m), ncol=m)
  gc()

  doSleep1 <- function(i) {
    library(pryr)
    fileConn<-file(paste("output_", Sys.getpid(), ".txt", sep=""))
    writeLines(paste("Memory in doSleep1 :", mem_used()), fileConn)
    close(fileConn)
    Sys.sleep(i);
  } # without, M and M2 will be serialized

  doSleep2 <- function(i) {
    library(pryr)
    fileConn<-file(paste("output_", Sys.getpid(), ".txt", sep=""))
    writeLines(paste("Memory in doSleep1 :", mem_used()), fileConn)
    close(fileConn)
    Sys.sleep(i);
  }
  environment(doSleep2) <- .GlobalEnv # without, M and M2 would be serialized

  doSleep3 <- function(i, M) { # M in parameter
    library(pryr)
    fileConn<-file(paste("output_", Sys.getpid(), ".txt", sep=""))
    writeLines(paste("Memory in doSleep1 :", mem_used()), fileConn)
    close(fileConn)
    Sys.sleep(i);
```

```

M[1,1]=0;
}
environment(doSleep3) <- .GlobalEnv # without, M is serialized but M2 too

cl <- makeCluster(2, type="PSOCK")
cat("doSleep1\n")
res = parLapply(cl, list(20,20), doSleep1)
stopCluster(cl)

cl <- makeCluster(2, type="PSOCK")
cat("doSleep2\n")
res = parLapply(cl, list(20,20), doSleep2)
stopCluster(cl)

cl <- makeCluster(2, type="PSOCK")
cat("doSleep3\n")
res = parLapply(cl, list(30,30), doSleep3, M)
stopCluster(cl)
NULL
}

parSleep()

```

2. reprendre la question précédente (dont la relance de R) en enlevant la/les ligne(s) `environment(doSleep...)` et comparer