R programming language: parallel computations

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Introductio

Loss distribution computation

Parallel computation R

R programming language: parallel computations

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Introduction

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Introduction

- A task to produce result Y out of input X may have steps that could be performed in parallel
- Example: you need to drink a 1L of beer
 - Ask a friend to help
 - The taks is, thus, parallelized among two throats

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Who does the job

- you can split the tasks in parallel steps, but who does the job?
- Nowadays, each PC usually has 2-8 CPU cores.
 - Each core can process a task!
- You can split the tasks across different computers
 - You can rent them at Microsoft, Amazon etc.

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How it works - MapReduce framework

- Map step assign to each work a task to perform
- Reduce step collect results from different workers and produce the overall result

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Basics

- Random numbers used for simulations are "pseudo-random", i.e. a sequence of number that looks random
- The starting point is called a "seed". Same seed same sequence
- Any simulation system must be reproducible

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Simulation

- assume we need to calculate 10 million of random numbers
- if we can split it among 4 CPU cores than each of them needs to generate only 2.5 millions
- we can assign to each core a different seed, generate the numbers and unite the numbers into one

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Implementation - I

Loss distribution - non-parallel

```
loss.dist <- function(seed, N)</pre>
  set.seed(seed)
  return(runif(N))
N < -4
seed <- 1
print(loss.dist(seed, N))
 [1] 0.2655087 0.3721239 0.5728534 0.9082078
```

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Implementation - II

• Explore your ressources

```
library(parallel)
print(detectCores())
[1] 4
```

Loss distribution - parallel

```
N <- 4
seed <- 1
print(loss.dist(seed, N))</pre>
```

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Using library parallel

- A lot of implementation details are hidden in R (e.g. compared to C#)
- General schema
 - Create a cluster (makeCluster)
 - Put in the scope of cluster objects (functions, variables etc) which are needed there (clusterExport)
 - Send a task (function together with input) to the cluster (parLapply) - similar to apply functions

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Mean calculation

- Take a trivial task as an example calculate sum of the sample
- divide N numbers sample on M workers
- calculate the sums for each workers (in parallel)
- collect the results and calculate their sum

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Sum parallel calculation - hands-on

```
nc <- 2 # number of cores
cl <- makeCluster(no_cores)</pre>
sq <- 1:8
L <- length(sq)
clusterExport(cl, list("sq", "nc", "L"))
# check first that the split is correct
parLapply(cl, 1:nc
  , function(x) sq[(1+(x-1)*L/nc):(x*L/nc)]
# [[1]]
  [1] 1 2 3 4
  [[2]]
  [1] 5 6 7 8
```

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Sum parallel calculation - hands-on | |

```
res <- parLapply(cl, 1:nc
  function(x) sum(sq[(1+(x-1)*L/nc):(x*L/nc)])
print(res)
  \lceil \lceil 1 \rceil \rceil
  Γ1] 10
  Γ[2]]
  Γ17 26
print(sum(unlist(res))) # collect results
# [1] 36
```

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Example - wrong implementation

• Farenheit to Celcius

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Task implement Farenheit to Celcius transform in parallel

 Take this as input (correct parLapply call is to be implemented)

```
library(parallel)
c <- function(t) t*5/9-32
nc <- 4
temps <- seq(10, 40, 10)
cl <- makeCluster(nc)
clusterExport(cl, list("temps"))</pre>
```

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Example - wrong implementation

- Example attempt parLapply(cl, 1:nc, function(x) c(temps[x]))
- However, temperatures are still in Farenheit, what is wrong here?
- Try to correct it

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Example - correction

```
library(parallel)
C \leftarrow function(t) t*5/9-32
nc <- 4
temps < seq(10, 40, 10)
cl <- makeCluster(nc)</pre>
clusterExport(cl, list("temps", "C"))
parLapply(cl, 1:nc, function(x) C(temps[x]))
\lceil \lceil 1 \rceil \rceil
[1] -26.44444
ΓΓ2]]
[1] -20.88889
[[3]]
[1] -15.33333
```

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Example - wrong implementation ||

- standard object c was overwritten and wasn't put in the scope of cluster
- Even worse it didn't throw an error since the cluster used the default object

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Pay attention

- Be careful with the scope of the cluster
- distribute carefully among work loaders

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Parallel loss distribution calculation

consider a simple loss generation

```
loss.dist <- function(seed, N)
{
   set.seed(seed)
   return(runif(N))
}
print(loss.dist(1,4))
# [1] 0.2655087 0.3721239 0.5728534 0.908207
[1] 0.2655087 0.3721239 0.5728534 0.9082078</pre>
```

• How to parallelize it?

```
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```

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```
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```

```
Parallel loss calculation

    Possible solution

 parallel.loss.dist <- function(seed, N)
    no_cores <- 2
    cl <- makeCluster(no cores)</pre>
    clusterExport(cl, list("loss.dist"))
    temp.res <- parLapply(
          сl
```

function(x) loss.dist(x, N = 3)

[1] 0.2655087 0.3721239 0-1848823 0.702374

, seed: (seed + 1)

return(unlist(temp.res))

print(parallel.loss.dist(1,4))

stopCluster(cl)

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Questions

- Why the first 2 numbers coincide with non-parallel version and the rest not?
- Where is the "Map" step and where is the "Reduce" step hidden in the code?

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Time measurement

Let's measure time with the following function (not optimal)

```
measure.time <- function(command)
  start.time <- Sys.time()
  eval(parse(text = command))
  end.time <- Sys.time()</pre>
  d <- difftime(end.time, start.time</pre>
         , units = "secs"))
  return(d)
 Example:
# cmd <- "sum(parallel.loss.dist(1, 1e+08))"</pre>
# measure.time(cmd)
```

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Time measurement ||

- play with N to see when it pays off to use parallel or non-parallel version
- adjust additionally the number of cores does it get faster?