#### Parallel R

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### Outline

- Motivation and introduction
- 2 snow
- 3 multicore
- 4 parallel
- 5 What else and references

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Why **R**? The **R** language has a lot of advantages:

- Open Source.
- Cross-Platform.
- Free.
- Many basic tools.
- R extensions.
- Arguably fast with R-way style and well-implemented R distribution.

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- Memory-bound: all data should fit in RAM.

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Solution: Parallel Execution. How exactly?

- Single-threaded: multiple CPU's (and cores).
- Memory-bound: spread data from one computer (master) to several computers (slaves).

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### Basic Terminology

The two types of parallelism:

- Implicit the OS abstracts parallelism from the user.
- Explicit user controls details of the process.

A computer cluster consists of a set of loosely or tightly connected computers that work together so that, in many respects, they can be viewed as a single system (Wiki).

Master/slave is a model of communication where one device or process has unidirectional control over one or more other devices (Wiki).

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#### Overview

#### snow

Explicit parallelism by using

Usage: clusters (works on Linux,

Windows, Mac OS X).

Solves: Single-threaded, memory-bound.

#### multicore [deprecated]

Implicit parallelism by using

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#### Hadoop

R + Hadoop, RHIPE, Segue.

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### **Epigraph**

"...R was not built in anticipation of the Big Data revolution.
R was born in 1995. Disk space was expensive, RAM even more so, and this thing called The Internet was just getting its legs. Notions of "large-scale data analysis" and "high-performance computing" were reasonably rare.
Outside of Wall Street firms and university research labs, there just wasn't that much data to crunch."

— Q. Ethan McCallum and Stephen Weston "Parallel R"

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### snow: quick look

Cool features:

General use case: Main word is cluster, provides explicit parallelism.

Examples: Monte Carlo simulations, bootstrapping, cross validation, ensemble machine learning algorithms.

Solves: Single-threaded, memory-bound.

 Different transport mechanisms between Master and Slaves: Sockets, MPI (rmpi), NWS (nws), PVM (rpvm).

• Good support of RNG (rsprng, rlecuyer).

Problems: No communication between the workers (slaves).

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Problems: No communication between the workers (slaves).

The input arguments must fit into memory when calling

Warning: snow function. Its up to the user to arrange high-performance

distributed file systems.

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#### Structure of API

```
Start and stop clusters: makeCluster, stopCluster.

Low-level (cluster-level) functions: cluster* — clusterApply, clusterApplyLB, clusterEvalQ, clusterCall, clusterSplit, etc.

High-level functions: par[L,S,A,R,C]apply — parallel versions of apply and related functions.

(Uniform) RNG: L'Ecuyer (package: rlecuyer), SPRNG [deprecated] (package: rsprng).

Timing: snow.time(expr) — very useful.
```

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### Start and stop clusters

#### Basic way:

First parameter is spec — specification. It is ssh-configurable unless you type "localhost":

```
cl <- makeCluster(c("localhost","localhost"), type = "SOCK")
stopCluster(cl)</pre>
```

Warning: Be aware of computational costs of cluster setup.

#### Low-level API

All of them are cluster\* and designed for computing *on* a cluster. Most interesting are as follows.

<pre>clusterApply(cl, x, fun,)</pre>	Jobs are being "recycled".
<pre>clusterApplyLB(cl, x, fun,)</pre>	Load Balancing version of clusterApply().
	Calls a function fun with identical
<pre>clusterCall(cl, fun,)</pre>	arguments on each node in the cluster cl and returns a list of the results.
<pre>clusterEvalQ(cl, expr)</pre>	Evaluates an expression expr on each node in the cluster cl;
	implemented using clusterCall().
<pre>clusterMap(cl, fun,,   MoreArgs = NULL, RECYCLE = TRUE)</pre>	Similar to mapply.

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### Example, K-means

#### Basic one-core way:

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# Before using snow it is easier to think \*apply-way:

```
results <- lapply(rep(25, 4),
  function(nstart)
    kmeans(Boston, 4, nstart=nstart)
)
i <- sapply(results,
  function(result)
    result$tot.withinss
)
result <- results[[which.min(i)]]</pre>
```

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```

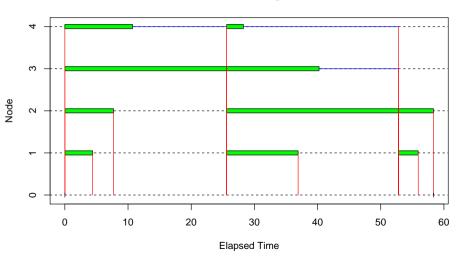
#### Finally snow version.

```
ignore <-
  clusterEvalQ(cl, {library(MASS); NULL})
results <-
  clusterApply(cl, rep(25, 4),
    function(nstart)
      kmeans(Boston, 4, nstart=nstart)
i <- sapply(results,
  function(result)
    result$tot.withinss
result <- results[[which.min(i)]]
```

clusterApply() uses a *robin-round* fashion for scheduling tasks for clusters. It means one time for every cluster. It could be not very wise to do that.

```
set.seed(123)
sleeptime <- abs(rnorm(10, 10, 10))
tm <- snow.time(clusterApply(cl, sleeptime, Sys.sleep))</pre>
```





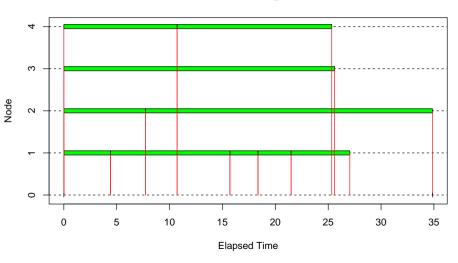
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So we waited for more than 50 seconds. A more efficient way would be pull the tasks to clusters when they are needed. This technique is called "load balancing". Function clusterApplyLB() uses that technique.

```
set.seed(123)
sleeptime <- abs(rnorm(10, 10, 10))
tm <- snow.time(clusterApplyLB(cl, sleeptime, Sys.sleep))</pre>
```





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So, here we waited for about 30 seconds. This is an improvement. The only wasted time was at the end.

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### High-level API

```
parLapply(cl, x, fun, ...) Parallel version of lapply().

parSapply(cl, X, FUN, ...,
    simplify = TRUE, USE.NAMES = TRUE)

parApply(cl, X, MARGIN, FUN, ...) Parallel version of apply().

parRapply(cl, x, fun, ...) Row apply() for matrix.

parCapply(cl, x, fun, ...) Column apply() for matrix.
```

The most useful is parLapply() function. It is different from clusterApply() because it splits the task into "equal" tasks.

```
parLapply

## function (cl, x, fun, ...)

## docall(c, clusterApply(cl, splitList(x, length(cl)), lapply,

## fun, ...))

## <environment: namespace:snow>
```

where splitList() is an internal snow function.

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parLapply() could be more efficient if you have more tasks than workers. Another situation — you send large arguments to parLapply(). Let's take a look at the example.

```
bigsleep <- function(sleeptime, mat) Sys.sleep(sleeptime)
bigmatrix <- matrix(0, 2000, 2000)
sleeptime <- rep(1, 100)</pre>
```

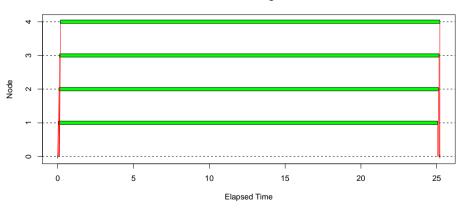
Firstly, let's try clusterApply().



Definitely not highly efficient. Those gaps are due to I/O time. Ideally we should have 25 seconds... Let's give parLapply() a try.

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# Load Balancing parLapply?

### Load Balancing parLapply?

Short answer: no, there is no such a function in snow package.

Good news: it is possible to write your own.

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#### Random Number Generation

#### There are 2 basic steps.

• Configure your cluster workers to use a generator.

```
library(rlecuyer)
clusterSetupRNG(cl, type = 'RNGstream')
## [1] "RNGstream"
```

2 Be happy to generate your numbers.

```
unlist(clusterCall(c1, runif, 1))
## [1] 0.12701112 0.75958186 0.72850979 0.09570262
```

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# Multicore [deprecated] : quick look

If it is deprecated, why even think about it?

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If it is deprecated, why even think about it? The reason is the package parallel. Wait a little bit...

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General use case: Main word is fork (thus no Windows support),

provides implicit parallelism.

Examples: lapply() runs for ages on your Intel Core i999.

Solves: Single-threaded.

No Windows support.

• No internal RNG support.

*Problems*: • Runs only on one computer.

Cannot be used with R GUI.

• No internal Load Balancing, however, it can be imitated.

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• Cannot be used with **R** GUI.

No internal Load Balancing, however, it can be imitated.

Warning: Jobs started by multicore share the same state (because of fork).

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We will only consider high-level API and let low-level to be out-of-scope.

mclapply()	Parallel version of lapply().
<pre>mcmapply()</pre>	Parallel version of mapply().
pvec()	Somewhat an high-level analog of
	low-level clusterSplit() function.
<pre>parallel() and collect()</pre>	<pre>parallel() creates a new process with fork(),</pre>
	evaluate expression in parallel and after that
	the result is retrieved by the collect().

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## Multicore: mclapply

```
mclapply() is a parallel lapply().
```

Syntax is as follows:

```
mclapply(X, FUN, ..., mc.preschedule = TRUE, mc.set.seed = TRUE,
    mc.silent = FALSE, mc.cores = getOption("mc.cores"))
where
```

- mc.preschedule = TRUE how jobs are created for X.
- mc.set.seed = TRUE do you need to randomly seed slaves, or fork it?
- mc.silent = FALSE hide info from 'stdout' for all parallel forked processes.
   'stderr' is not affected.
- o mc.cores == getOption("mc.cores") number of workers (not cores, actually) to start.

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#### Multicore: mclapply, mc.preschedule

### Meaning:

- TRUE: divide data in mc.cores-jobs beforehand and fork it to mc.cores-processes.
- FALSE: for each piece of data construct a new job (up to mc.cores).

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#### Rule of thumb: use

- TRUE: you don't need load balance (for instance, if there are lot's of values in the data).
- FALSE: the variance of job completion is very high (so, you need load balance).

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## parallel

General use case: Main word is mainstream,

almost a wrapper of snow and multicore packages.

Examples: Anything above.

Solves: Single-threaded and (partially) memory-bound.

Preinstalled into R since 2.14.0.

 Full RNG support with no dependency on rlecuyer package.

Cool features:

Almost nothing to learn (if you are still awake).

Can be easily used on any platform including Windows.

Highly compatible with snow and multicore.

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Almost nothing to learn (if you are still awake).

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Warning:

On Windows you can't use more than one machine. It also can be difficult to configure multiple Linux machines.

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## parallel : detectCores()

How many cores?

```
library(parallel)
mc.cores <- detectCores()
mc.cores
## [1] 8</pre>
```

Warning: It is important to take into account that you maybe have hyper-threading.

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## parallel RNG

Unlike snow package no additional packages (like rlecuyer) are needed.

Fork (no Windows) way:

```
RNGkind("L'Ecuyer-CMRG")
unlist(mclapply(rep(1,4), runif))
## [1] 0.3768615 0.3824588 0.3845725 0.9092709
```

### Cluster way:

```
detach("package:snow", character.only=TRUE)
library(parallel)
RNGkind("L'Ecuyer-CMRG")
cl <- makeCluster(2, type="PSOCK")
unlist(clusterCall(cl, function(x) runif(2)))
## [1] 0.3114024 0.9506436 0.6032429 0.8057068
stopCluster(cl)</pre>
```

## parallel RNG: reproducible results

Basic way to get reproducible results would be mc.reset.stream() — the parallel random number generator is reinitialized using the current seed on the master.

```
detach("package:snow", character.only=TRUE)
library(parallel)
RNGkind("L'Ecuyer-CMRG")
cl <- makeCluster(2, type="PSOCK")</pre>
clusterSetRNGStream(cl, 123)
unlist(clusterCall(cl. function(x) runif(2)))
   [1] 0.1663742 0.3898457 0.3411064 0.9712727
clusterSetRNGStream(cl. 123)
unlist(clusterCall(cl. function(x) runif(2)))
## [1] 0.1663742 0.3898457 0.3411064 0.9712727
stopCluster(cl)
```

## Differences from multicore and snow

Let's sum up the differences between modern parallel package and his predecessors.

parallel > multicore

- Prefix mc in mcfork(), mcexit(), mckill(), mcparallel(), mccollect(), mc.cores.
- Different default value of mc.cores argument.
- New mc.reset.stream() function.

Also useful detectCores() is added.

 $parallel \neq snow$ 

- New function clusterSetRNGStream() initializes parallel RNG.
- snow.time() function not included.
- makeCluster() supports additional types FORK.

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## Out of scope

We covered 3 (2.5 really) packages: snow, multicore, parallel. What else?

- Revolution Analytics for each package for iteration over a set of values.
- MapReduce via Java Hadoop: RHIPE (negotiator between you with your MapReduce functions and Hadoop).
- Segue for Amazon Elastic MapReduce lovers. Be aware of terminating clusters.
- doRedis.
- http://cloudNumbers.com
- R and GPUs: gputools etc.

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### Literature

#### Main reference:

Data Analysis in the Distributed World



O'REILLY®

O. Etban McCallum & Stephen Weston

#### **Useful links:**

- Advanced R by Hadley Wickham.
- The R Inferno by Patrick Burns.
- R Packages by Hadley Wickham.
- Writing R Extensions.
- Los Angeles R Users Group: Parallelization in R, Revisited by Ryan R. Rosario.
- Package parallel manual.

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