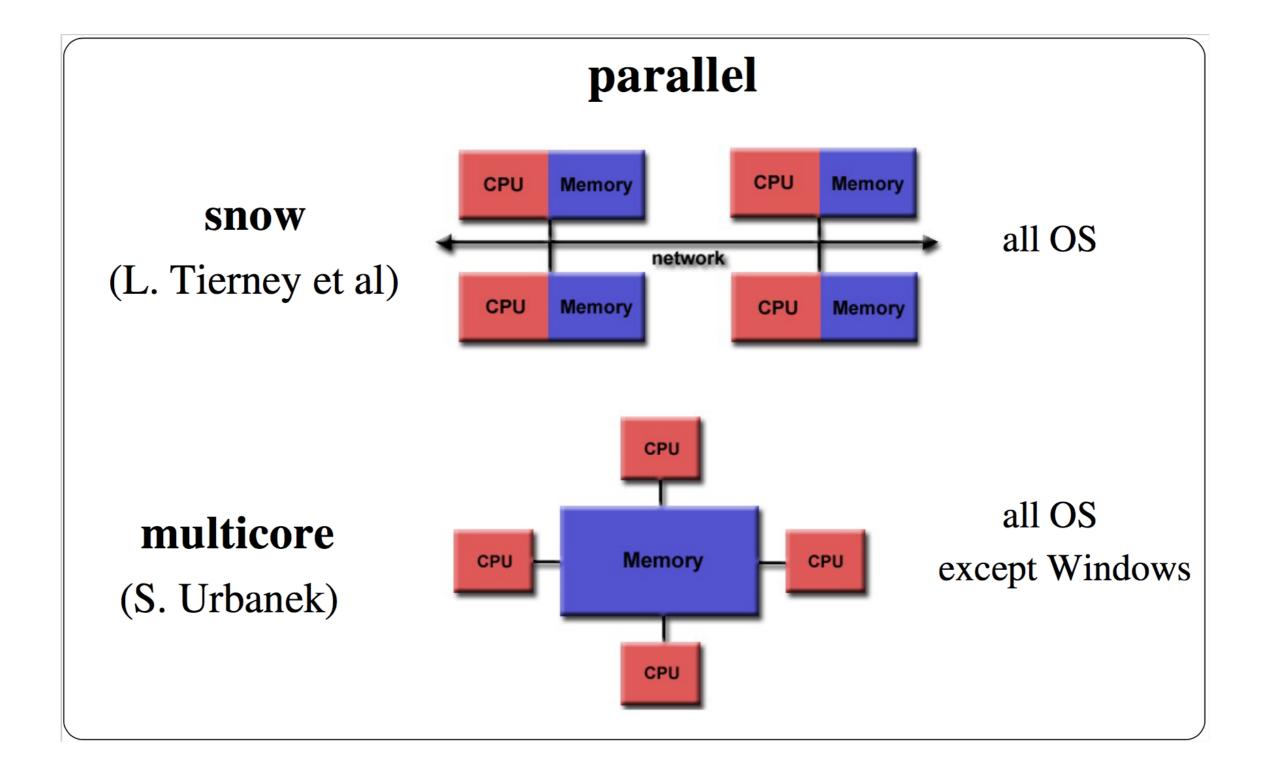
Cluster basics

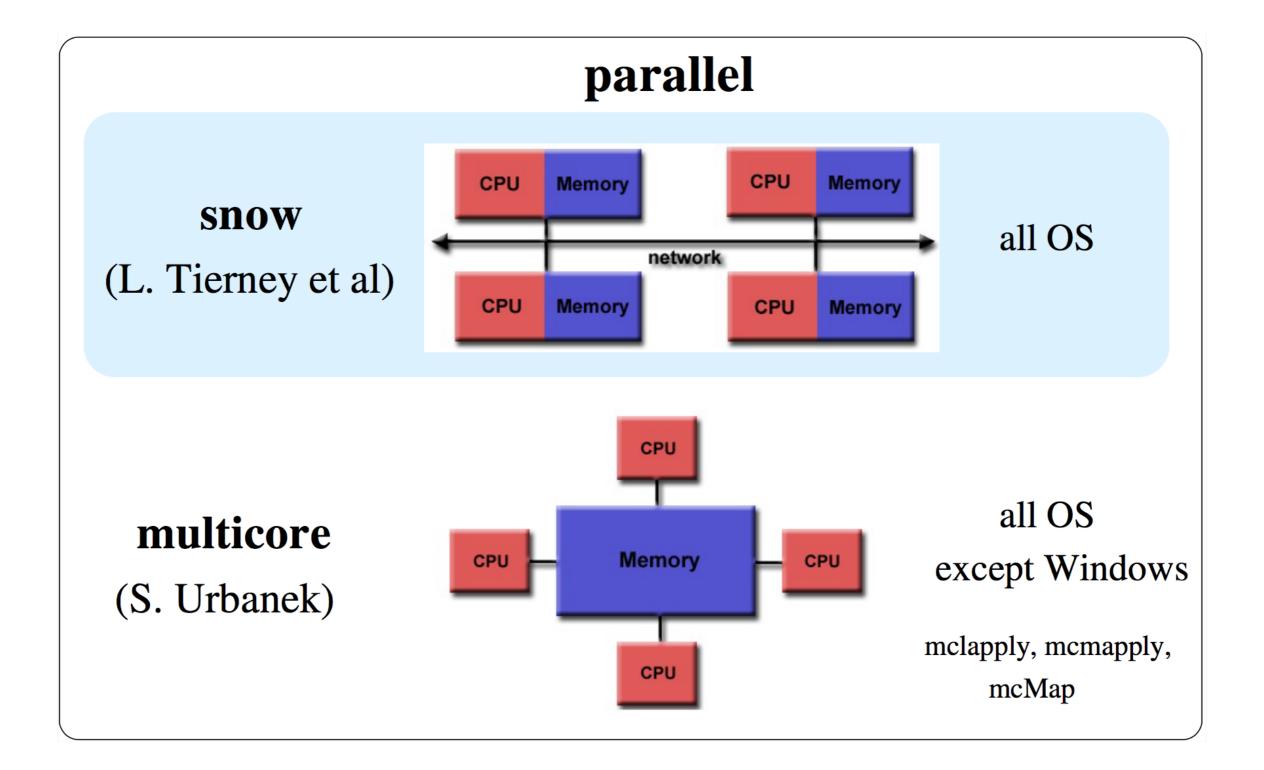
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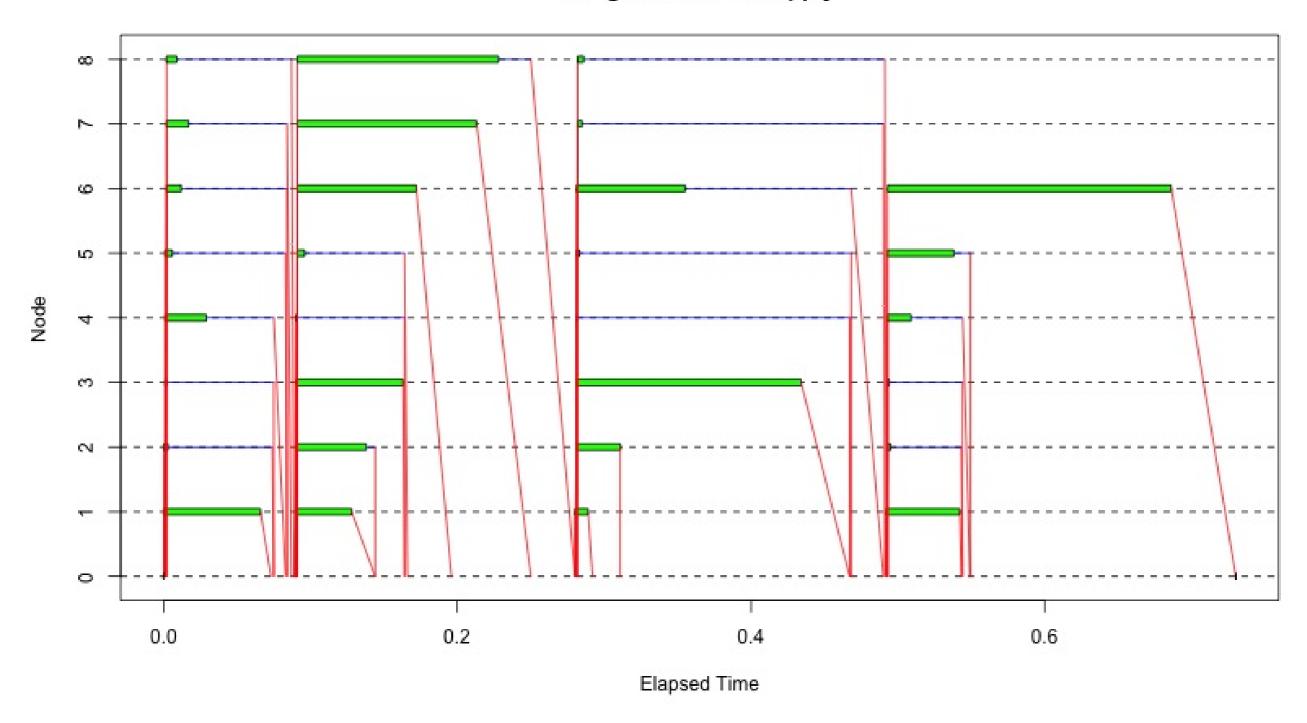
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Usage with clusterApply





Supported backends

Socket communication (default, all OS platforms)

```
cl <- makeCluster(ncores, type = "PSOCK")</pre>
```

• Workers start with an empty environment (i.e. new R process).

Supported backends

Forking (not available for Windows)

```
cl <- makeCluster(ncores, type = "FORK")</pre>
```

Workers are complete copies of the master process.

Supported backends

Using the MPI library (uses Rmpi)

```
cl <- makeCluster(ncores, type = "MPI")</pre>
```

Let's practice!

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The core of parallel

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Core functions

Main processing functions:

- clusterApply()
- clusterApplyLB()

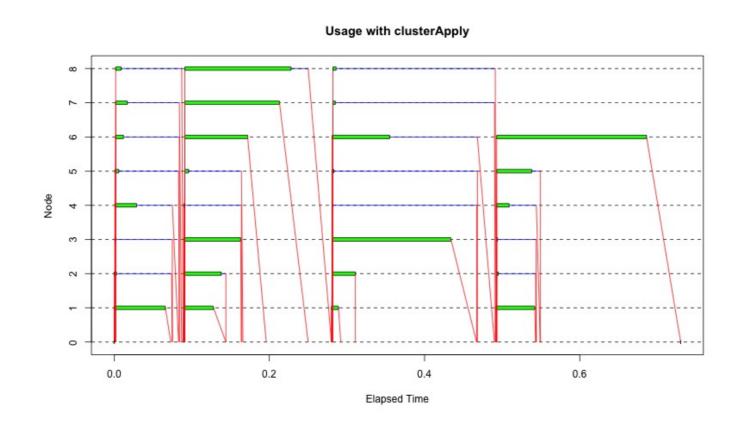
Wrappers:

- parApply(), parLapply(), parSapply()
- parRapply(), parCapply()
- parLapplyLB(), parSapplyLB()

clusterApply(): Number of tasks

```
clusterApply(cl, x = arg.sequence, fun = myfunc)
```

length(arg.sequence) = number of tasks (green bars)





Parallel vs. sequential

Not all embarrassingly parallel applications are suited for parallel processing.

Processing overhead:

- Starting/stopping cluster
- Number of messages sent between nodes and master
- Size of messages (sending big data is expensive)

Things to consider:

- How big is a single task (green bar)
- How much data need to be sent
- How much gain is there by running it in parallel → benchmark

Let's practice!

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Initialization of nodes

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Why initialize?

- Each cluster node starts with an empty environment (no libraries loaded).
- Repeated communication with the master is expensive.
- Example:

```
clusterApply(cl, rep(1000, n), rnorm, sd = 1:1000)
```

- Master sends a vector of 1:1000 to all n tasks (n can be very large).
- Good practice: Master initializes workers at the beginning with everything that stays constant or/and is time consuming. Examples:
 - Sending static data
 - Loading libraries
 - Evaluating global functions

clusterCall()

• Evaluates the same function with the same arguments on all nodes.

Example:

```
cl <- makeCluster(2)
clusterCall(cl, function() library(janeaustenr))
clusterCall(cl, function(i) emma[i], 20)

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

```
[1] "She was the youngest of the two daughters of a most affectionate,"
[2]]
[1] "She was the youngest of the two daughters of a most affectionate,"
```

clusterEvalQ()

• Evaluates a literal expression on all nodes (equivalent to evalq())

Example:

```
[[1]]
[1] "Sense & Sensibility" "Pride & Prejudice" "Mansfield Park"

[[2]]
[1] "Sense & Sensibility" "Pride & Prejudice" "Mansfield Park"
```

clusterExport()

Exports given objects from master to workers.

Example:

```
books <- get_books()
cl <- makeCluster(2)
clusterExport(cl, "books")
clusterCall(cl, function() print(books))</pre>
```

Let's practice!

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Subsetting data

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Data chunks

- Each task applied to different data (data chunk)
- Data chunks are passed to workers as follows:
 - 1. Random numbers generated on the fly
 - 2. Passing chunks of data as argument
 - 3. Chunking on workers' side

Data chunk as random numbers

```
myfunc <- function(n, ...) mean(rnorm(n, ...))
clusterApply(cl, rep(1000, 20), myfunc, sd = 6)</pre>
```

Data chunk as argument (1)

- Dataset is chunked into several blocks on master
- Each block passed to worker via an argument
- Incorporated into higher level functions (parApply() etc)

```
cl <- makeCluster(4)
mat <- matrix(rnorm(12), ncol=4)</pre>
```

```
[,1] [,2] [,3] [,4]

[1,] 1.1540263 -2.180922 0.5322614 0.5578128

[2,] -1.8763588 -1.625226 0.4058091 -0.5532732

[3,] -0.1685597 -1.089104 0.1770636 0.5483025
```

Data chunk as argument (2)

```
Sum of columns (colSums(mat)):
```

```
parCapply(cl, mat, sum)
unlist(clusterApply(cl, as.data.frame(mat), sum))
```

• Sends each worker a column of mat

Chunking on workers' end

Example of matrix multiplication $M \times M$:

```
n <- 100
M <- matrix(rnorm(n * n), ncol = n)
clusterExport(cl, "M")

mult_row <- function(id) apply(M, 2, function(col) sum(M[id,] * col))

clusterApply(cl, 1:n, mult_row) %>% do.call(rbind, .)
```

Let's practice!

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