Tackling Big Data with R

New features and old concepts for handling large and streaming data in practice

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R Foundation

Overview

- Motivation
- Custom connections
- Data processing pipelines
- Parallel processing
- Back-end experiments: Hadoop, RDFS
- Call for participation

Motivation

- R's in memory model is fast
 - RAM prices declining steadily (unlike CPUs), [ca. \$8/Gb for server RAM now]
 - Billion+ rows in R workable
- Problem 1: parallelization
 - s <- split(df, ...) ### slow and ineffcient!</pre>
 - y <- mclapply(s, function(x) ...)</pre>
 - splitting up data is expensive
- Problem 2: streaming
 - conceptually cannot have all data at once

Old, simple idea: chunking

- Process data in (big) chunks
- Parallelization:
 - feed each process/worker with chunks, collect results
 - can process chunks in parallel (if the processing can be independent); no copying
- Streaming:
 - keep a mutable state
 - process chunks as they come in, modifying state and creating results
- Issue: R has no explicit framework/API for this

Connections

- R has connections: abstraction for data access and transport - completely back-end opaque!
- New in R 3.0.0: custom connection support
 - packages can create new connection implementations
 - some examples:
 - zmqc OMQ PUB/SUB connections read from OMQ streaming feeds directly
 - hdfsc read files from HDFS just like any other file

```
f = HDFS("/data/foo")
d = read.table(f)
```

Data pipeline

mean(read.table(HDFS("foo"))\$x)

Source - delivers data

```
connection (text or binary)
```

Data parser - converts data format to R objects

data frame (or other R-native object)

Filtering, processing, computing, ...

result (aggregates, models, graphics, ...)

Streaming

```
Source - delivers data
connection (text or binary)
Data parser - converts data format to R objects
```

```
data frame
(or other R-native object)
```

```
Filtering, processing, computing, ...
```

result

(aggregates, models, graphics, ...)



Parallel processing



Proposal: Chunks in a pipeline

- Connections
 - define available classes of data sources contribute!
- Read from sources in big chunks
- Parsers
 - transform data representation to R objects contribute!
- Compute
 - algorithms that work on chunks contribute!
 (serial processing + mutable state = streaming, independence = parallel)
- Collect
 - algorithms to combine parallel chunks

contribute!

Example: streaming

 Use OMQ PUB/SUB: buffered per subscriber (slow subscribers don't affect others; can detect dropped

```
recor

• Read

• Upda

• Upda

• Serve

Rhttp

Rhttp

recor

feed = zmqc("ipc:///my-feed.0mq", "r")

max = 1000

state = numeric()

while (TRUE) {

    d = read.table(feed, FALSE, nrows=max)

    mix = c(state * 0.9, table(d[,2]))

    state = tapply(mix, names(mix), sum)

    if (any(state <= 1)) state = state[state > 1]

}
```

Parallel processing

- At least three stages:
 - split (often implicit)
 - compute
 - combine
- Define functions using this paradigm simple examples:
- cc.sum <- function(x) cc(x, sum, sum)</pre>

cc.table <- function(x) cc(x, table, function(x) tapply(x, names(x), sum))</pre>

Practical considerations

- The implementation can be seamless: use special "distributed vector" class and dispatch on it
- Typically source is big, so splitting is implicit since the data does not reside in R (e.g. sequence in a file)
- Leverage distributed storage: run computing where the chunks are stored

Examples:

- Hadoop
- RDFS

Hadoop

- A lot of companies invest in Hadoop clusters (we have to live with it even if there are many better solutions)
- Literal map/reduce based on key/value is very inefficient for R since it is not a vector operation
- Hadoop can be (ab)used for chunk-wise processing: streaming mode - use HDFS chunks as input, compute is map on the entire chunk, combine is reduce

Example

• Aggregate point locations by ZIP code (match points against ZCTA US/Census 2010 shapefiles)

```
r <- read.table(hmr(
    hinput("/data/2013/06"),
    function(x)
       table(zcta2010.db()[
            inside(zcta2010.shp(), x[,4], x[,5]), 1]),
    function(x) ctapply(x, names(x), sum)))
```

- Fairly native R programming
- Implicit defaults (read.table parser, conversion of named vectors to key/value entries)
- Result is an HDFS connection

R Distributed File System - Experiment

- Purely R-based (R client, R server, R code)
- Uses Rserve for fast access (no setup cost, optional authentication, users switching, transport encryption for free)
- Any storage available (RData, ASCII, ...), all storage is R-native parsing step can be removed
- No name node, all nodes are equal
- Scales only to moderate cluster sizes (hundreds of nodes), but is very fast (milliseconds for job setup, no need to leave R)

Call for Participation

- More users, more use cases
 - is this powerful enough?
 - if not, what is missing?
- Make it part of R
 - so developers can rely on it
- Start writing functions and packages
 - help to create critical mass
- Theoretical work
 - methods and approaches that give bounds for approximation error, necessary assumptions etc.

Related work

- Purdue Univ: Divide/Recombine
 - results for linear model approximations
 - RHipe very specialized vehicle for the above using specific version and brand of Hadoop
- Iterators (also used by foreach)
 - idea of running code in iterations; does include chunks
 - focused on inner code (chunk processing)

Conclusions

- New in R 3.0.0: custom connections, to be used as building blocks for data pipelines
- Read from connections in chunks, compute and collect
- Generic framework that can be applied to streaming and parallel processing
- Let us work together to see if it is powerful enough to build an official R interface that everyone can use and contribute to
- Back-end agnostic testing on Hadoop and RDFS

Contact

- Most packages available on **RForge.net** (source also on GitHub)
 - http://RForge.net/zmqc
 - http://RForge.net/hdfsc
- Remaining packages (iotools, rdfs, ...) in the process of being pushed, check RForge.net and
 - https://github.com/s-u

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