Benchmarking multiple out-of-memory strategies

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1 Introduction

In many large-data situations, it is impractical to load and retain data in R's working memory space. We have had a look at HDF5, SQLite and tabix-indexed text as possible solutions to problems arising with memory constraints. We'll call these "out-of-memory" (OOM) approaches

How can we obtain data on which approach will be most effective for a given task? Comparative benchmarking is a very useful skill and we give a very rudimentary account of this here.

2 The harness

It is common to speak of a program that drives other programs as a "harness" (see wikipedia for related discussion). We have such a program in ph525x:

```
library(bench00M)
bench00M
## function (NR = 5000, NC = 100, times = 5, inseed = 1234, methods = list(.h5RoundTrip,
##
       .ffRoundTrip, .slRoundTrip, .dtRoundTrip, .bmRoundTrip))
## {
       nel = NR * NC
##
##
       set.seed(inseed)
       x = array(rnorm(nel), dim = c(NR, NC))
##
##
       cbind(NR = NR, NC = NC, times = times, do.call(rbind, lapply(methods,
##
           function(z) getStats(times, x, rtfun = z))), units = "microsec")
## }
## <environment: namespace:bench00M>
```

This program is going to help us assess performance of various OOM approaches. We consider a very limited problem, that of managing data that could reside in an R matrix. The main parameters are

- NR and NC: row and column dimensions
- times: number of benchmark replications for averaging
- inseed: a seed for random number generation to ensure reproducibility
- methods: a list of methods

The methods parameter is most complex. Each element of the list is assumed to be a function with the matrix to be managed via OOM as the first argument, some additional parameters, and a parameter intimes that gives the number of benchmark replicates.

Our objective is to produce a table that looks like

```
> b1
    NR NC times
                      meth
                                         ingFull
                                                      ing1K
1 5000 100
                                       9.4100810 14.2984402
              5
                      hdf5 10.71714
              5
2 5000 100
                        ff 25.34365 63.0977338 4.4320688
3 5000 100
                    sqlite 174.89003 105.1254638 28.4717496
              5
4 5000 100
              5 data.table 49.35190
                                       7.9871552 13.9007588
5 5000 100
              5 bigmemory 23.39697
                                       0.9660878 0.9950034
```

where each method listed in meth is asked to perform the same task a fixed number of times for averaging. The construction of the table occurs by binding together metadata about the task and method to the result of getStats. We'll leave the details of getStats to independent investigation.

3 An example method (OOM benchmarker for HDF5 matrix)

Let's look at the method for HDF5:

```
benchOOM:::.h5RoundTrip
## function (x, chunkIn = c(1000, 10), inLevel = 0, intimes = 1)
## {
##
       if (file.exists("ex_hdf5file.h5"))
##
           file.remove("ex hdf5file.h5")
##
       requireNamespace("rhdf5")
       h5createFile("ex_hdf5file.h5")
##
##
       h5createDataset("ex_hdf5file.h5", "x", c(nrow(x), ncol(x)),
##
           storage.mode = "double", chunk = chunkIn, level = inLevel)
##
       mw = microbenchmark(h5write(x, "ex_hdf5file.h5", name = "x"),
##
           times = intimes)
       mr = microbenchmark(h5read("ex_hdf5file.h5", name = "x"),
##
##
           times = intimes)
       msel = microbenchmark(ysel <- h5read("ex_hdf5file.h5", name = "x",</pre>
##
           index = list(4001:5000, 1:100)), times = intimes)
##
       stopifnot(all.equal(ysel, x[4001:5000, ]))
##
       list(mwrite = mw, ingFull = mr, ing1K = msel, times = intimes,
##
           method = "hdf5")
##
## }
## <environment: namespace:bench00M>
```

The program has three main phases

- HDF5-related setup, cleaning out any previous archives and establishing the basic target file
- Benchmarking of data export via h5write
- Benchmarking of ingestion via h5read with various restrictions

The results of microbenchmark are assembled in a list.