Benchmarks for Discrete Fourier Transforms in R

Andrew J. Barbour

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Abstract

The base DFT calculator in R, stats::fft, uses the Mixed-Radix algorithm of Singleton (1969). In this vignette we show how this calculator compares to FFT in the fftw package (Krey et al., 2011), which uses the FFTW algorithm of Frigo and Johnson (2005). For univariate DFT computations, the methods are nearly equivalent with two exceptions which are not mutually exclusive: (A) the series to be transformed is very long, and especially (B) when the series length is not highly composite. In both exceptions the algorithm FFT outperforms fft.

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1 Benchmarking function

We use both functions in their default state, and ask them to transform the same univariate random series. Benchmark information comes from the rbenchmark program, and the versatile plyr and reshape2 packages are used to manipulate the information for this presentation; ggplot2 is used for plotting. First we load the libraries needed:

```
rm(list = ls())
library(fftw)
library(rbenchmark)
library(plyr)
library(reshape2)
library(ggplot2)
```

and create a benchmark function:

```
reps <- 10
dftbm <- function(nd, repls = reps) {
    set.seed(1234)
    x <- rnorm(nd, mean = 0, sd = 1)
    bmd <- benchmark(replications = repls, fftw::FFT(x), stats::fft(x))
    bmd$num_dat <- nd
    bmd$relative[is.na(bmd$relative)] <- 1  # NA happens.
    return(bmd)
}</pre>
```

2 Highly composite (HC) series

It's well known that DFT algorithms are most efficient for "Highly Composite Numbers", specifically multiples of (2,3,5).

So, we create a vector of series lengths we wish to benchmark

```
(nterms.even \leftarrow round(2^seq.int(from = 4, to = 20, by = 1)))
##
    [1]
                       32
                                 64
                                         128
                                                  256
                                                           512
                                                                   1024
                                                                            2048
              16
    [9]
            4096
                             16384
                                      32768
                                                                262144
##
                     8192
                                               65536
                                                       131072
                                                                          524288
## [17] 1048576
```

¹ This is the reason for the stats::nextn function.

and use it with lapply and the benchmark function previously defined. These data are further distilled into a usable format with ldply:

```
bench.even <- function() {
    benchdat.e <- plyr::ldply(lapply(X = nterms.even, FUN = dftbm))
}
bench.even()</pre>
```

In order to plot the results, we need to perform some map/reduce operations on the data (Wickham, 2011):

```
pltbench <- function(lentyp=c("even", "odd")){</pre>
  benchdat <- switch(match.arg(lentyp), even=benchdat.e, odd=benchdat.o)</pre>
  stopifnot(exists("benchdat"))
  tests <- unique(benchdat$test)</pre>
  ## subset only information we care about
  allbench.df.drp <- subset(benchdat,</pre>
        select=c(test, num_dat, user.self, sys.self, elapsed, relative))
  ## reduce data.frame with melt
  allbench.df.mlt <- reshape2::melt(allbench.df.drp,</pre>
                                      id.vars=c("test", "num_dat"))
  ## calculate the summary information to be plotted:
  tmpd <- plyr::ddply(allbench.df.mlt,</pre>
                       .(variable, num_dat),
                       summarise,
                       summary="medians", # just a name
                       value=ggplot2::mean_cl_normal(value)[1,1])
  ## create copies for each test and map to data.frame
  allmeds <<- plyr::ldply(lapply(X=tests,</pre>
                                   FUN=function(x,df=tmpd){
                                         df$test <- x; return(df)</pre>
                                       }))
  ## plot the benchmark data
  g <- ggplot(data=allbench.df.mlt,
               aes(x=log10(num_dat),
                   y=log2(value),
                   # 1/sqrt(n) standard errors [assumes N(0,1)]
                   ymin=log2(value*(1-1/sqrt(reps))),
```

```
ymax=log2(value*(1+1/sqrt(reps))),
                colour=test,
                group=test)) +
     scale_colour_discrete(guide="none") +
     theme_bw()+
     ylim(c(-11,11))+
     xlim(c(0.5,6.5))+
     ggtitle(sprintf("DFT benchmarks of %s length series",toupper(lentyp)))
## add previous summary curves if exist
if (exists("allmeds.prev")){
   g <- g + geom_path(size=1.5, colour="dark grey", data=allmeds.prev,
                       aes(group=test))
## create a facetted version
g2 <- g + facet_grid(variable~test) #, scales="free_y")</pre>
## add the summary data as a line
g3 <- g2 + geom_path(colour="black", data=allmeds, aes(group=test))
## and finally the data
print(g4 <<- g3 + geom_pointrange())</pre>
```

For each row of the figure we plot summary curves² so we can easily inter-compare the benchmark data.

3 Non highly composite (NHC) series

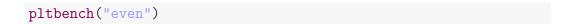
DFT algorithms can have drastically reduced performance if the series length is not highly composite (NHC). We now test NHC series by adding one to the HC series-length vector (also restricting the total length for sanity's sake):

```
nterms.odd <- nterms.even + 1
nterms.odd <- nterms.odd[nterms.odd < 50000] # painfully long otherwise!</pre>
```

and performing the full set of benchmarks again:

² Based on this post:

http://geokook.wordpress.com/2012/12/29/row-wise-summary-curves-in-faceted-ggplot2-figu



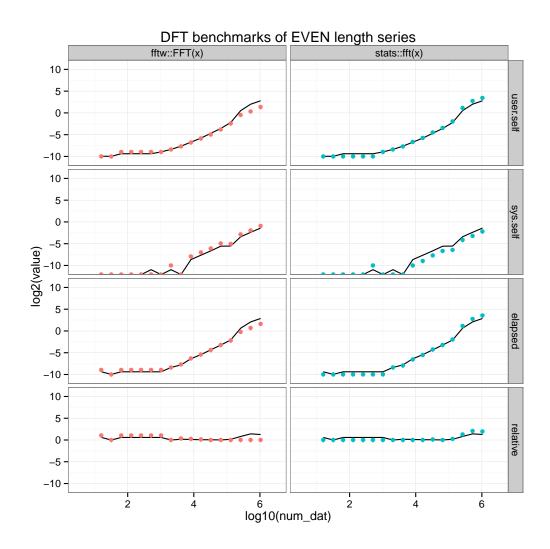


Figure 1: DFT benchmark results for HC series lengths.

```
bench.odd <- function() {
    benchdat.o <- plyr::ldply(lapply(X = nterms.odd, FUN = dftbm))
}</pre>
```

bench.odd() # FAIR WARNING: this can take a while!!

We can now visualize the results, with the addition of the HC summary curves:

4 Conclusion

Figures 2 and 3 compare the DFT calculations for HC and NHC length series. For univariate DFT computations, the methods are nearly equivalent with two exceptions which are not mutually exclusive: (A) the series to be transformed is very long, and especially (B) when the series length is not highly composite. In both exceptions the algorithm FFT outperforms fft. In the case of exception (B), both methods have drastically increased computation times; hence, zero padding should be done to ensure the length does not adversely affect the efficiency of the DFT calculator.

References

- Frigo, M. and Johnson, S. G. (2005). The design and implementation of FFTW3. *Proceedings of the IEEE*, 93(2):216–231. Special issue on "Program Generation, Optimization, and Platform Adaptation".
- Krey, S., Ligges, U., and Mersmann, O. (2011). fftw: Fast FFT and DCT based on FFTW. R package version 1.0-3.
- Singleton, R. C. (1969). An Algorithm for Computing the Mixed Radix Fast Fourier Transform. *IEEE Transactions on Audio and Electroacoustics*, AU-17(2):93–103.
- Wickham, H. (2011). The split-apply-combine strategy for data analysis. Journal of Statistical Software, 40(1):1–29.

allmeds.prev <- allmeds
pltbench("odd")</pre>

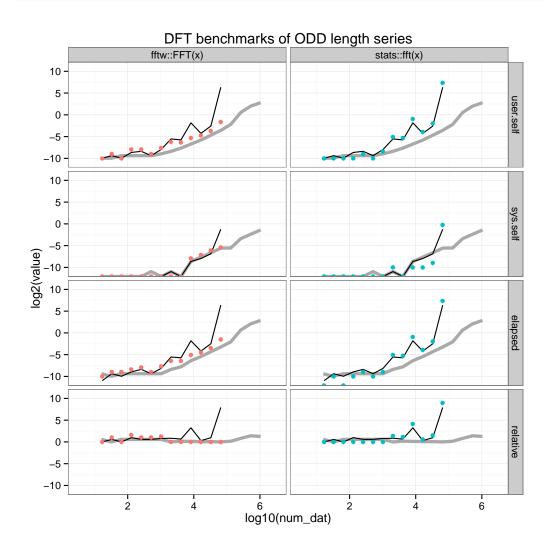


Figure 2: DFT benchmark results for NHC series lengths. We also show the summary curves for the HC results to highlight the drastic degradation in performance.