# Dip Test Distributions, P-values, and other Explorations

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#### Abstract

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Keywords: MPFR, Abitrary Precision, Multiple Precision Floating-Point, R.

- [1] "/tmp/diptest.Rcheck"
- [2] "/sfs/u/maechler/R/library/64-linux-MM-only"
- [3] "/sfs/s/linux/rhel3\_amd64/app/R/R\_local/library\_F19"
- [4] "/sfs/s/linux/rhel3\_amd64/app/R/Bioconductor/library\_2.13"
- [5] "/sfs/u/maechler/R/library/64-linux"
- [6] "/sfs/u/maechler/R/D/r-devel/F17-64-inst/library"

## 1. Introduction

FIXME: Need notation

 $D_n := dip( runif(n) );$ 

but more generally,

$$D_n(F) := D(X_1, X_2, \dots, X_n), \quad \text{where } X_i \text{ i.i.d. } , X_i \sim F.$$

Hartigan and Hartigan (1985) in their "seminal" paper on the dip statistic  $D_n$  already proved that  $\sqrt{n} D_n$  converges in distribution, i.e.,

$$\lim_{n \to \infty} \sqrt{n} \ D_n \stackrel{\mathcal{D}}{=} \ D_{\infty}. \tag{2}$$

A considerable part of this paper is devoted to explore the distribution of  $D_{\infty}$ .

## 2. History of the diptest R package

Hartigan (1985) published an implementation in Fortran of a concrete algorithm, where the code was also made available on Statlib<sup>1</sup>

<sup>&</sup>lt;sup>1</sup>Statlib is now a website, of course, http://lib.stat.cmu.edu/, but then was *the* preferred way for distributing algorithms for statistical computing, available years before the existence of the WWW, and entailing e-mail and (anonymous) FTP

On July 28, 1994, Dario Ringach, then at NY University, asked on Snews (the mailing list for S and S-plus users) about distributions and was helped by me and then about dyn.load problems, again helped by me. Subsequently he provided me with S-plus code which interfaced to (a f2ced version of) Hartigan's Fortran code, for computing the dip statistic. and ended the (then private) e-mail with

I am not going to have time to set this up for submission to StatLib. If you want to do it, please go ahead.

Regards, Dario

- several important bug fixes; last one Oct./Nov. 2003

However, the Fortran code file http://lib.stat.cmu.edu/apstat/217, was last changed Thu 04 Aug 2005 03:43:28 PM CEST.

We have some results of the dip.dist of before the bug fix; notably the "dip of the dip" probabilities have changed considerably!!

- see rcs log of ../../src/dip.c

# 3. 21st Century Improvement of Hartigan<sup>2</sup>'s Table

((

Use listing package (or so to more or less "cut & paste" the nice code in ../../stuff/new-simul.Rout-1e6

))

# 4. The Dip in the Dip's Distribution

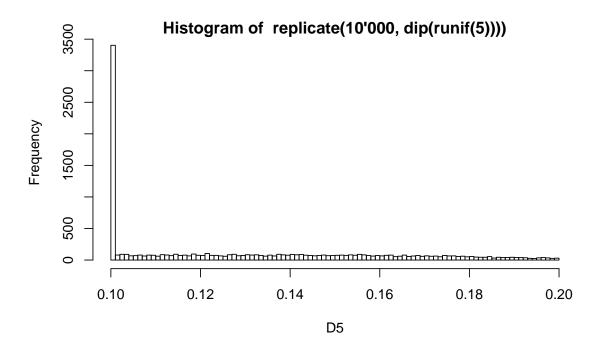
We have found empirically that the dip distribution itself starts with a "dip". Specifically, the minimal possible value of  $D_n$  is  $\frac{1}{2n}$  and the probability of reaching that value,

$$P\left[D_n = \frac{1}{2n}\right],\tag{3}$$

is large for small n.

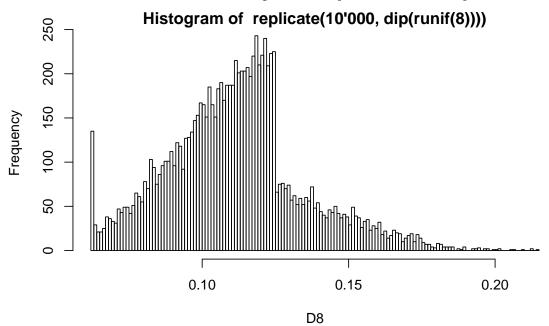
E.g., consider an approximation of the dip distribution for n=5,

```
R> require("diptest") # after installing it ..
R> D5 <- replicate(10000, dip(runif(5)))
R> hist(D5, breaks=128, main = "Histogram of replicate(10'000, dip(runif(5)))")
```



which looks as if there was a bug in the software — but that look is misleading! Note how the phenomenon is still visible for n = 8,

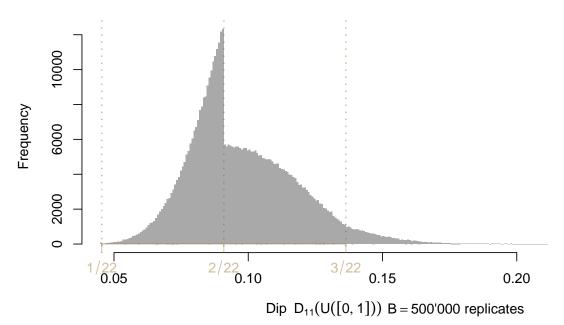
R> D8 <- replicate(10000, dip(runif(8)))
R> hist(D8, breaks=128, main = "Histogram of replicate(10'000, dip(runif(8))))")



Note that there is another phenomenon, in addition to the point mass at 1/(2n), particularly visible, if we use many replicates,

R> set.seed(11)
R> n <- 11</pre>

R> B.s11 <- 500000
R> D11 <- replicate(B.s11, dip(runif(n)))</pre>



FIXME:

use '../../stuff/sim-minProb.R' and '../../stuff/minProb-anal.R'

Further, it can be seen that the *maximal* dip statistic is  $\frac{1}{4} = 0.25$  and this upper bound can be reached simply (for even n) using the data  $(0,0,\ldots,0,\ 1,1,\ldots,1)$ , a bi-point mass with equal mass at both points.

# 5. P-values for the Dip Test

Note that it is not obvious how to compute P-values for "the dip test", as that means determining the distribution of the test statistic, i.e.,  $D_n$  under the null hypothesis, but a natural null,  $H_o: F \in \{F \text{cadlag} \mid f := \frac{d}{dx}Fisunimodal\}$  is too large. Hartigans'(1985) argued for using the uniform U[0,1] i.e.,  $F'(x) = f(x) = \mathbf{1}_{[0,1]}(x) = [0 \le x \le 1]$  (Iverson bracket) instead, even though they showed that it is not quite the "least favorable" one. Following Hartigans', we will define the P-value of an observed  $d_n$  as

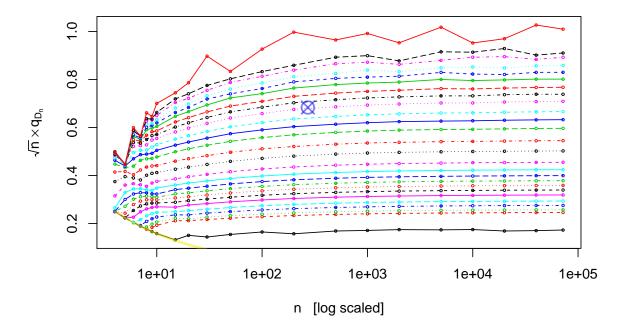
$$P_{d_n} := P[D_n \ge d_n] := P[\operatorname{dip}(U_1, \dots, U_n) \ge d_n], \text{ where } U_i \sim U[0, 1], \text{ i.i.d.}$$
 (4)

### 5.1. Interpolating the Dip Table

Because of the asymptotic distribution,  $\lim_{n\to\infty} \sqrt{n} D_n \stackrel{\mathcal{D}}{=} D_{\infty}$ , it is makes sense to consider the " $\sqrt{n}D_n$ "-scale, even for finite n values:

R> data(qDiptab)
R> dnqd <- dimnames(qDiptab)
R> (nn. <- as.integer(dnqd[["n"]]))</pre>

```
[1]
                                              10
                                                                30
                                                                       50
                                                    15
                                                          20
                                    5000 10000 20000 40000 72000
[12]
       100
             200
                   500
                        1000
                              2000
R> matplot(nn., qDiptab*sqrt(nn.), type ="o", pch=1, cex = 0.4,
         log="x", xlab="n
                             [log scaled]",
         ylab = expression(sqrt(n) %*% q[D[n]]))
R> ## Note that 1/2n is the first possible value (with finite mass),,
R> ## clearly visible for (very) small n:
R> lines(nn., sqrt(nn.)/(2*nn.), col=adjustcolor("yellow2",0.5), lwd=3)
R> P.p <- as.numeric(print(noquote(dnqd[["Pr"]])))</pre>
             0.01
                      0.02
                              0.05
                                                       0.3
                                                               0.4
 [9] 0.5
             0.6
                      0.7
                              0.8
                                      0.9
                                               0.95
                                                       0.98
                                                               0.99
[17] 0.995
             0.998
                      0.999
                              0.9995
                                      0.9998
                                               0.9999
                                                       0.99995 0.99998
[25] 0.99999 1
R> ## Now look at one well known data set:
R > D <- dip(x <- faithful waiting)
R > n <- length(x)
R> points(n, sqrt(n)*D, pch=13, cex=2, col= adjustcolor("blue2",.5), lwd=2)
R> ## a simulated (approximate) P-value for D is
R> mean(D <= replicate(10000, dip(runif(n)))) ## ~ 0.002</pre>
[1] 0.0021
```



but we can use our table to compute a deterministic (but still approximate, as the table is from simulation too) P-value:

```
R> ## We are in this interval:
R> n0 <- nn.[i.n <- findInterval(n, nn.)]
R> n1 <- nn.[i.n +1] ; c(n0, n1)
[1] 200 500
R> f.n <- (n - n0)/(n1 - n0)# in [0, 1]
R> ## Now "find" y-interval:
```

[1] 0.001809527

R> ## 0.018095

Finally, in May 2011, after several year of people asking for it, I have implemented a dip.test function which makes use of a — somewhat more sophisticated — interpolation scheme like the one above, to compute a P-value. As qDiptab has been based on  $10^6$  samples, the interpolation yields accurate P-values, unless in very extreme cases. Here is the small (n=63) example from Hartigan<sup>2</sup>,

where, from a P-value of 8.7%, we'd conclude that there is not enough evidence against unimodality.

## 5.2. Asymptotic Dip Distribution

We have conducted extensive simulations in order to explore the limit distribution of  $D_{\infty}$ , i.e., the limit of  $\sqrt{n} D_n$ , (2).

Our current R code is in '../../stuff/asymp-distrib.R ' but the simulation results (7 Megabytes for each n) cannot be assumed to be part of the package, nor maybe even to be simply accessible via the internet.

## 6. Less Conservative Dip Testing

#### 7. Session Info

R> toLatex(sessionInfo())

- R Under development (unstable) (2014-11-22 r67037), x86\_64-unknown-linux-gnu
- Locale: LC\_CTYPE=C, LC\_NUMERIC=C, LC\_TIME=en\_US.UTF-8, LC\_COLLATE=C, LC\_MONETARY=en\_US.UTF-8, LC\_MESSAGES=C, LC\_PAPER=en\_US.UTF-8, LC\_NAME=C, LC\_ADDRESS=C, LC\_TELEPHONE=C, LC\_MEASUREMENT=en\_US.UTF-8, LC\_IDENTIFICATION=C
- Base packages: base, datasets, grDevices, graphics, methods, stats, tools, utils

• Other packages: diptest 0.75-6

## References

Hartigan JA, Hartigan PM (1985). "The Dip Test of Unimodality." *Annals of Statistics*, **13**, 70–84.

Hartigan PM (1985). "Computation of the Dip Statistic to Test for Unimodality." *Applied Statistics*, **34**, 320–325.

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