

In this discussion, we look at confidence intervals in the Wilcoxon Sign-Rank Test for one sample. The data comes from an article by Jeffries, Voris and Yang, “Diversity and Distribution of the Pedunculate Barnacles ... on the Scillarid Lobster,” *Crustaceana*, 1984, as quoted by Mendenhall, Beaver & Beaver, *Introduction to Probability sand Statistics*, 14th ed., Brooks Cole, 2013. The carapace lengths (in mm) of randomly selected lobsters caught in the seas near Singapore are simulated consistent with the given data.

We assume that the sample X_1, \dots, X_N comes from a continuous distribution that is symmetric about its mean μ . The Wilcoxon signed rank statistic is computed as follows. To test

$$\begin{aligned} \mathcal{H}_0 &: \mu = \mu_0; \\ \mathcal{H}_a &: \mu \neq \mu_0. \end{aligned}$$

we order the values $|X_i - \mu_0|$ from lowest to highest. after throwing out any zero values, we have n nonzero terms. We rank them 1 to n . S_+ is the sum of the ranks corresponding to terms for which $X_i - \mu_0$ is positive. The null hypothesis is rejected at the significance level α if $S_+ \geq c$ or $S_+ \leq \frac{n(n+1)}{2} - c$ where $2P(S_+ \geq c) = \alpha$.

Theorem 1. Fix μ_0 . Assume that there are no ties and no zeros among the values $|X_i - \mu_0|$ for $i = 1 \dots, n$. Then S_+ is equal to the number of pairs (i, j) such that $i \leq j$ and $X_i + X_j \geq 2\mu_0$.

Proof. Observe that the number of such pairs is unchanged if the observations are rearranged. Thus we may suppose that the observations are arranged from smallest to largest $X_{(1)} < \dots < X_{(n)}$. Let r_i denote the rank of $|X_{(i)} - \mu_0|$. Let k be the index where the $X_{(i)} - \mu_0$ change sign, that is

$$X_{(k)} - \mu_0 < 0 < X_{(k+1)} - \mu_0$$

Then the ranks decrease to k and then increase from $k + 1$:

$$r_1 > r_2 > \dots > r_k, \quad r_{k+1} < r_{k+2} < \dots < r_n.$$

For example if the sample is 12.2, 10.0, 11.1, 15.5, 14.4 then $X_{(1)} = 10.0$, $X_{(2)} = 11.1$, $X_{(3)} = 12.2$, $X_{(4)} = 14.4$ and $X_{(5)} = 17.7$ so if $\mu_0 = 13.9$ then $|X_{(1)} - \mu_0| = 3.9$, $|X_{(2)} - \mu_0| = 2.8$, $|X_{(3)} - \mu_0| = 1.7$, $|X_{(4)} - \mu_0| = .5$ and $|X_{(5)} - \mu_0| = 3.8$ thus $r_1 = 5$, $r_2 = 3$, $r_3 = 2$, $r_4 = 1$ and $r_5 = 4$, and finally $k = 3$ and $S_+ = r_{k+1} + \dots + r_n = 1 + 4 = 5$.

Let $\chi_{\mathcal{P}}$ be the indicator function which equals 1 if \mathcal{P} is true and 0 otherwise. Then the number of pairs satisfying the condition is

$$\sum_{i \leq j} \chi\{X_i + X_j \geq 2\mu_0\} = \sum_{i \leq j} \chi\{X_{(i)} + X_{(j)} \geq 2\mu_0\} \tag{1}$$

$$= \sum_{j=k+1}^n \sum_{i=1}^j \chi\{X_{(j)} - \mu_0 \geq \mu_0 - X_{(i)}\} \tag{2}$$

$$= \sum_{j=k+1}^n \sum_{i=1}^n \chi\{r_i \leq r_j\} \tag{3}$$

$$= \sum_{j=k+1}^n r_j \tag{4}$$

$$= S_+ \tag{5}$$

(1) holds because the number of pairs does not depend on the order of the variables. (2) holds because if $i \leq j \leq k$ then $X_{(i)} - \mu_0 < X_{(j)} - \mu_0 < 0$ so that $X_{(i)} + X_{(j)} < 2\mu_0$ and does not add to the sum. $k + 1 \leq i \leq j$ if and only if $r_i \leq r_j$ and $j < i$ if and only if $r_j < r_i$. On the other hand if $i \leq k$ then $r_i \leq r_j$ if and only if $\mu_0 - X_{(i)} = |X_{(i)} - \mu_0| \leq |X_{(j)} - \mu_0| = X_{(j)} - \mu_0$ so $X_{(i)} + X_{(j)} \geq 2\mu_0$. On the other hand, $r_i \leq r_j$ if and only if $\mu_0 - X_{(i)} = |X_{(i)} - \mu_0| > |X_{(j)} - \mu_0| = X_{(j)} - \mu_0$ so $X_{(i)} + X_{(j)} < 2\mu_0$. It means that for a given j and any i , $X_{(i)} + X_{(j)} \geq 2\mu_0$ if and only if $r_i < r_j$ so we may replace the sum in (3). Now there are exactly r_j i s such that $r_i \leq r_j$ giving (4). But (5) is the definition of S_+ . \square

The confidence interval associated to the Sign-Rank test \mathcal{H}_0 above is the following: we consider the set of values

$$\mathcal{A} = \left\{ \frac{X_i + X_j}{2} : 1 \leq i \leq j \leq n \right\}$$

If there are no ties, then S_+ is the number of elements of \mathcal{A} that are not smaller than μ_0 . If we sort \mathcal{A} into $\{A_{(1)} \leq A_{(2)} \leq \dots \leq A_{(m)}\}$ where $m = \frac{n(n+1)}{2}$.

The one and two-sided critical values for S_+ may be computed as follow. Assuming \mathcal{H}_0 , the chance that the sign of the i th observation $X_i - \mu_0$ is positive or negative is equally likely. Thus the distribution of S_+ is obtained by looking at the histogram of values

$$\pm 1 \pm 2 \pm 3 \pm \dots \pm n$$

each of which has an equal chance 2^{-n} of occurring. They range from 0 to m . The pmf $p(x)$ of S_+ is symmetrical about the mean $\mu_{S_+} = \frac{n(n+1)}{4}$. The one-sided upper critical value is $P(S_+ \geq c_1) = \sum_{i \geq c_1} p(i) = \alpha$. The two sided critical value c satisfies $P(S_+ \geq c) = P(S_+ \leq m - c) = \frac{\alpha}{2}$. These numbers are tabulated in tables A 13 and A 15 of the text. We show how to compute the values of these tables in the code, albeit inefficiently.

Then the two-sided confidence interval for μ at the level α where c is the two-sided critical value that satisfies $P(S_+ \geq c) = \frac{\alpha}{2}$ is given by

$$(A_{(m+1-c)}, A_{(c)}).$$

Data Set Used in this Analysis :

```
# Math 3080                Carapace Data                April 19, 2014
# Treibergs
#
# From an article by Jeffries, Voris and Yang, "Diversity and Distribution
# of the Pedunculate Barnacles on the Scillarid Lobster," Crustaceana,
# 1984 as quoted by Mendenhall, Beaver & Beaver, Introduction to
# Probability sand Statistics, 14th ed., Brooks Cole, 2013.
#
# The carapace lengths (in mm) of randomly selected lobsters caught in
# the seas near Singapore are simulated consistent with the given data.
Length
79.2
81.1
56.4
78.2
64.8
64.4
69.7
69.8
45.3
64.3
65.9
69.4
70.0
68.6
52.6
63.6
54.5
60.6
```

R Session:

```
R version 2.13.1 (2011-07-08)
Copyright (C) 2011 The R Foundation for Statistical Computing
ISBN 3-900051-07-0
Platform: i386-apple-darwin9.8.0/i386 (32-bit)
```

```
R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.
```

```
Natural language support but running in an English locale
```

```
R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.
```

```
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
```

Type 'q()' to quit R.

[R.app GUI 1.41 (5874) i386-apple-darwin9.8.0]

[History restored from /Users/andrejstreibergs/.Rapp.history]

```
>
> tt=read.table("M3082DataCarapace.txt",header=T)
> attach(tt)
> tt
  Length

> t2=read.table("M3082DataCarapace.txt",header=T)
> attach(t2)
  Length
1    79.2
2    81.1
3    56.4
4    78.2
5    64.8
6    64.4
7    69.7
8    69.8
9    45.3
10   64.3
11   65.9
12   69.4
13   70.0
14   68.6
15   52.6
16   63.6
17   54.5
18   60.6
> table(Length)
Length
45.3 52.6 54.5 56.4 60.6 63.6 64.3 64.4 64.8 65.9 68.6 69.4 69.7 69.8  70
   1   1   1   1   1   1   1   1   1   1   1   1   1   1   1
78.2 79.2 81.1
   1   1   1
```

```

> ##### CRITICAL VALUES FOR SIGNED-RANK TEST #####
> ##### ILLUSTRATE DETAILS FOR N=5 #####
> di=5
> #### MATRIX WHOSE COLS ARE ALL POSSIBLE POSITIVE RANK SUMS #####
> M=matrix(1:(di*2^di),ncol=2^di)
> for(i in 1:di){M[i,]=rep(i*0:1,each=2^(i-1),length=2^di)}
> M
      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11] [,12] [,13]
[1,]    0    1    0    1    0    1    0    1    0    1    0    1    0
[2,]    0    0    2    2    0    0    2    2    0    0    2    2    0
[3,]    0    0    0    0    3    3    3    3    0    0    0    0    3
[4,]    0    0    0    0    0    0    0    0    4    4    4    4    4
[5,]    0    0    0    0    0    0    0    0    0    0    0    0    0
      [,14] [,15] [,16] [,17] [,18] [,19] [,20] [,21] [,22] [,23] [,24] [,25]
[1,]     1     0     1     0     1     0     1     0     1     0     1     0
[2,]     0     2     2     0     0     2     2     0     0     2     2     0
[3,]     3     3     3     0     0     0     0     3     3     3     3     0
[4,]     4     4     4     0     0     0     0     0     0     0     0     4
[5,]     0     0     0     5     5     5     5     5     5     5     5     5
      [,26] [,27] [,28] [,29] [,30] [,31] [,32]
[1,]     1     0     1     0     1     0     1
[2,]     0     2     2     0     0     2     2
[3,]     0     0     0     3     3     3     3
[4,]     4     4     4     4     4     4     4
[5,]     5     5     5     5     5     5     5
> ##### PDF OF S+ #####
> p=table(margin.table(M,2))/2^di;sum(p)
[1] 1
> p
      0      1      2      3      4      5      6      7      8
0.03125 0.03125 0.03125 0.06250 0.06250 0.09375 0.09375 0.09375 0.09375
      9     10     11     12     13     14     15
0.09375 0.09375 0.06250 0.06250 0.03125 0.03125 0.03125

# ##### WORK OUT TAIL PROBABILITIES #####
> cump=cumsum(p)
> np=length(p);np;di*(di+1)/2+1
[1] 16
[1] 16
> names(cump)=np-1:np
> #### N=5 ONE-TAILED CRITICAL FOR SIGNED-RANK TEST (TABLE A13) ####
> #### c1 AND ALPHA TABLE #####
> cump
      15      14      13      12      11      10      9      8      7
0.03125 0.06250 0.09375 0.15625 0.21875 0.31250 0.40625 0.50000 0.59375
      6      5      4      3      2      1      0
0.68750 0.78125 0.84375 0.90625 0.93750 0.96875 1.00000

```

```

> cum2p=1-2*cumsum(p)
> names(cum2p)=np-1:np
> ##### N=5 TWO-TAILED CRITICAL FOR SIGNED-RANK TEST (TABLE A15) #####
> ##### c AND ALPHA TABLE #####
> cum2p[1:(np/2)]
      15      14      13      12      11      10      9      8
0.9375 0.8750 0.8125 0.6875 0.5625 0.3750 0.1875 0.0000
>
> ##### COMPUTE TWO-SIDED CRITICAL VALUES FOR CARAPACE DATA #####

> di=length(Length); di
[1] 18
> M=matrix(1:(di*2^di),ncol=2^di)
> for(i in 1:di){M[i,]=rep(i*0:1,each=2^(i-1),length=2^di)}
> p=table(margin.table(M,2))/2^di;sum(p)
[1] 1
> cum2p=1-2*cumsum(p)
> np=length(p);np;di*(di+1)/2+1
[1] 172
[1] 172
> names(cum2p)=np-1:np
> ##### N=18 TWO-TAILED CRITICAL FOR SIGNED-RANK TEST (TABLE A15) ###
> ##### c AND ALPHA TABLE #####
> cum2p[1:(np/2)]
      171      170      169      168      167      166      165
0.99999237 0.99998474 0.99997711 0.99996185 0.99994659 0.99992371 0.99989319
      164      163      162      161      160      159      158
0.99985504 0.99980927 0.99974823 0.99967194 0.99958038 0.99946594 0.99932861
      157      156      155      154      153      152      151
0.99916077 0.99895477 0.99871063 0.99842072 0.99806976 0.99766541 0.99719238
      150      149      148      147      146      145      144
0.99663544 0.99599457 0.99525452 0.99440002 0.99342346 0.99230957 0.99103546
      143      142      141      140      139      138      137
0.98959351 0.98796844 0.98612976 0.98406982 0.98176575 0.97918701 0.97632599
      136      135      134      133      132      131      130
0.97315216 0.96963501 0.96576691 0.96150970 0.95683289 0.95172119 0.94614410
      129      128      127      126      125      124      123
0.94006348 0.93346405 0.92631531 0.91857147 0.91023254 0.90126038 0.89161682
      122      121      120      119      118      117      116
0.88129425 0.87026215 0.85848236 0.84595490 0.83264923 0.81853485 0.80361176
      115      114      113      112      111      110      109
0.78785706 0.77124786 0.75379181 0.73547363 0.71627045 0.69620514 0.67527008
      108      107      106      105      104      103      102
0.65345001 0.63078308 0.60726166 0.58288574 0.55770111 0.53170776 0.50492096
      101      100      99      98      97      96      95
0.47738647 0.44912720 0.42015839 0.39054108 0.36030579 0.32947540 0.29811859
      94      93      92      91      90      89      88
0.26627350 0.23397064 0.20129395 0.16827393 0.13495636 0.10142517 0.06771851
      87      86
0.03388214 0.00000000

```

```

>
> Length
[1] 79.2 81.1 56.4 78.2 64.8 64.4 69.7 69.8 45.3 64.3 65.9 69.4 70.0 68.6 52.6
[16] 63.6 54.5 60.6
> ##### GENERATE ALL (X[I]+X[J])/2 FOR I \le J #####
> xPx=1:(n*(n+1)/2)
> n=length(Length);n
[1] 18
> for(j in 1:n){for(i in 1:j){xPx[j*(j-1)/2+i]=(Length[i]+Length[j])/2}}
> sxPx=sort(xPx)
> ##### SORTED VALUES (X[I]+X[J])/2 FOR I \le J #####
> sxPx
[1] 45.30 48.95 49.90 50.85 52.60 52.95 53.55 54.45 54.50 54.50 54.80 54.85
[13] 55.05 55.45 55.60 56.40 56.60 56.95 57.35 57.50 57.55 57.55 57.65 58.10
[25] 58.45 58.50 58.50 58.70 59.05 59.25 59.40 59.45 59.65 60.00 60.20 60.35
[37] 60.40 60.60 60.60 60.60 61.00 61.15 61.15 61.20 61.30 61.55 61.75 61.95
[49] 62.10 62.10 62.15 62.25 62.25 62.45 62.50 62.50 62.70 62.90 63.05 63.10
[61] 63.20 63.20 63.25 63.60 63.95 64.00 64.20 64.30 64.35 64.40 64.55 64.60
[73] 64.60 64.75 64.80 65.00 65.10 65.15 65.15 65.20 65.30 65.35 65.40 65.90
[85] 65.90 66.10 66.35 66.45 66.50 66.50 66.65 66.70 66.70 66.80 66.85 66.85
[97] 66.85 66.90 67.00 67.05 67.05 67.10 67.10 67.15 67.20 67.25 67.25 67.30
[109] 67.30 67.40 67.65 67.80 67.80 67.80 67.85 67.95 68.60 68.75 69.00 69.15
[121] 69.20 69.30 69.40 69.40 69.55 69.60 69.70 69.70 69.75 69.80 69.85 69.90
[133] 69.90 70.00 70.85 70.90 71.25 71.30 71.40 71.50 71.75 71.80 72.00 72.05
[145] 72.35 72.55 72.70 72.75 72.95 73.40 73.50 73.80 73.90 73.95 74.00 74.10
[157] 74.30 74.45 74.50 74.60 74.85 75.25 75.40 75.45 75.55 78.20 78.70 79.20
[169] 79.65 80.15 81.10

> ##### TWO-SIDED CI ON MU FOR CARAPACE DATA #####
> ##### ALPHA = .05 #####
> c05=131;n1=di*(di+1)/2-c05+1;c(sxPx[n1],sxPx[c05])
[1] 61.00 69.85
> ##### CANNED CI FOR MU IN SIGNED-RANK TEST #####
> wilcox.test(Length,conf.int=T)

```

Wilcoxon signed rank test

```

data: Length
V = 171, p-value = 7.629e-06
alternative hypothesis: true location is not equal to 0
95 percent confidence interval:
 61.00 69.85
sample estimates:
(pseudo)median
 66.1

```

```

> ##### ALPHA = .01 #####
> c01=143;n1=di*(di+1)/2-c01+1;c(sxPx[n1],sxPx[c01])
[1] 59.05 72.00

> wilcox.test(Length,conf.int=T,conf.level=.99)

Wilcoxon signed rank test

data: Length
V = 171, p-value = 7.629e-06
alternative hypothesis: true location is not equal to 0
99 percent confidence interval:
 58.70 72.05
sample estimates:
(pseudo)median
      66.1

> ### CANNED CI DOES NOT USE C=143 OF TABLE A15 FOR 99.0 % TEST #####
> c01=144;n1=di*(di+1)/2-c01+1;c(sxPx[n1],sxPx[c01])
[1] 58.70 72.05

```


Histogram of Length

