## Mann-Whitney-Wilcoxon test: rationale

Assume that we have two samples  $\mathbf{X}=(x_1,...,x_{n1})$  and  $\mathbf{Y}=(y_1,...,y_{n2})$  and we want determine if they come from the same generating distribution, that is we want to test  $H_0:L_x=L_y$  using a nonparametric test, i.e. without assuming one parametric distribution, e.g. Normal, Binomial... Furthermore, assume that  $L_x$  and  $L_y$  are both continuous distributions and that  $x_i$  is independent of  $y_j$  for all i,j. One possible solution: the **Mann-Whitney-Wilcoxon** test otherwise known as the Wilcoxon's rank sum test.

Note: alternative versions of the Mann-Whitney-Wilcoxon test exists in case we specify the null hypothesis differently.

Let's uncover the mathematics behind this nonparametric test and its normal approximation. In a second time, we will perform the test and some simulations in the R language.

#### Basic notation and test statistic

Let  $S=(x_1,...,x_{n1},y_1,...,y_{n2})$  denote the pooled sample and R=rank(S), that is the ordered rank of all observation. Let us denote  $N=n_1+n_2$ . Finally, let us consider the r.v.  $R_X$ , which is the sum of the ranks of the observations from our first sample  $x_1,...,x_{n1}$  in the pooled sample S, defined as

$$R_X = \sum_{i=1}^{n1} R_i$$

The Mann-Whitney-Wilcoxon test statistic is usually given by

$$W = 2R_X - n_1(N+1)$$

We are only concerned with testing  $H_0: L_x = L_y$ .

## Distribution of the test statistic and large-sample approximation

The distribution W depends on the sample sizes  $n_1$  and  $n_2$ . If we do not have tables which gives us a critical value to test  $H_0$ , we can estimate it using simulations. If  $n_1$  and  $n_2$  are large enough, there is a possibility of a 'large-sample' approximation.

**Asymptotic approximation:** Using the CLT, we can prove that, for larges values of  $n_1$  and  $n_2$ , W has the following distribution

$$W \sim N\left(0, \ n_1 n_2 \frac{N+1}{3}\right)$$

We can use the function pnorm() in R to compute an approximate p-value.

## Working example

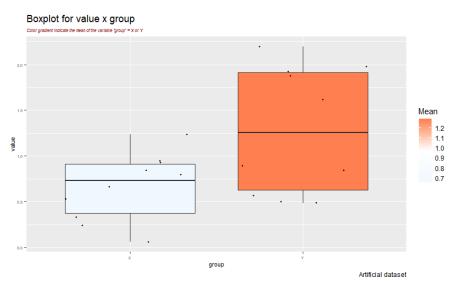
**Example:** The following values are uniform samples with  $\mathbf{X} \sim U_{[0,2]}$  and  $\mathbf{Y} \sim U_{[0,2,2,2]}$  of sample sizes  $n_1 = n_2 = 10$ 

$$X = 0.93322789, \ 0.67038191, \ 0.32563512, \ 0.79224003, \ 0.06078346, \ 0.24176974, \ 0.85233131, \ 1.23571576, \ 0.52641652, \ 0.95264774$$

$$Y = 1.9246380, 0.4977573, 0.5608602, 2.1985466, 1.8834824, 0.4852981, 0.8896968, 1.9792509, 0.8396966, 1.6180317$$

Using the exact distribution of W, then the asymptotic approximation, what is the conclusion of the test at a level of significance of 5% testing  $H_0: L_x = L_y$ , the data is coming from the same generating process or equivalently the temperatures are similar in both cities.

## Visualizing the data



# Mann-Whitney-Wilcoxon test using inbuilt function in R

```
1 # 1. Mann-Whitney-Wilcoxon using inbuilt function in R
2
3 set.seed(2023)
4 X = runif(50, min = 0, max = 2)
5 Y = runif(50, min = 0.2, max = 2.2)
6
7 wilcox.test(X,Y)
8 # Wilcoxon rank sum exact test
9 #
10 # data: X and Y
11 # W = 27, p-value = 0.08921
12 # alternative hypothesis: true location shift is not equal to 0
```

Conclusion: we can NOT reject  $H_0$ : 'both data come from the same generating distribution' at a significance level of 5% since for those samples, p-value>0.05.

## Asymptotic Mann-Whitney-Wilcoxon test in R

```
1 # 2. Asymptotic Mann-Whitney-Wilcoxon using inbuilt function in R
2 set. seed (2023)
3 X = runif(10, min = 0, max = 2)
4 Y = runif(10, min = 0.2, max = 2.2)
5 S = c(X, Y)
6 R = rank(S)
7 \text{ Rx} = \text{sum}(R[1:length(X)])
8 W = 2*Rx + (12*(N+1))
9 \text{ varW} = 50*50*((N+1)/3)
10
11 pnorm(W, mean = 0, sd = sqrt(varW), lower.tail = FALSE)
12 # 0.0008313872
13
14 # Conclusion: using the asysmptotic approximation, we reject HO: Lx = Lv that
        the both
15 # data come from the same generating distribution.
```

Conclusion: we can reject  $H_0$ : 'both data come from the same generating distribution' at a significance level of 5% since for those samples, p-value < 0.05.

# Expectation of the sum of ranks and sum or squared ranks

We know that, under  $H_0: L_x = L_y$ , the  $R_i$ 's are uniformly distributed on the set  $\{1,...,N\}$ . When computing the expectation, it involves the sum of the first N integers, given by the formula  $\frac{N(N+1)}{2}$ . It can be proven by induction (appendix 1).

So the expectation of the ranks and the squared ranks are respectively given by

$$E[R_i] = \frac{1}{N} \sum_{i=1}^{N} R_i = \frac{1}{N} \frac{N(N+1)}{2} = \frac{N+1}{2}$$

$$E[R_i^2] = \frac{1}{N} \sum_{i=1}^N R_i^2 = \frac{1}{N} \frac{N(N+1)(2N+1)}{6} = \frac{(N+1)(2N+1)}{6}$$

#### Variance of the ranks

By definition, the variance of the random variable  $R_i$  is given by

$$var(R_i) = E[R_i^2] - (E[R_i])^2$$

Then replacing by the results that we derived on the previous slide, we get

$$var(R_i) = \frac{(N+1)(2N+1)}{6} - \left(\frac{N+1}{2}\right)^2 = \frac{N^2 - 1}{12}$$

Since  $R_i$  is not independent from  $R_j$ , then the variance of the sum is equal to the sum of the variances plus a covariance term.

## Covariance term among ranks

We first note that  $\sum_{i=1}^{N} R_i$  is a constant and therefore its variance is 0.

$$\underbrace{\operatorname{var}(\sum_{i=1}^{N} R_i)}_{0} = \underbrace{\sum_{i=1}^{N} var(R_i)}_{\frac{N(N^2-1)}{12}} + \underbrace{\sum_{i=1}^{N} \sum_{j=1}^{N-1} cov(R_i, R_j)}_{N(N-1)cov(R_i, R_j)} \qquad \text{for } i \neq j.$$

So we have that

$$-\frac{N(N^2 - 1)}{12} = N(N - 1)cov(R_i, R_j)$$
$$-\frac{N(N - 1)(N + 1)}{12} = N(N - 1)cov(R_i, R_j)$$
$$-\frac{(N + 1)}{12} = cov(R_i, R_j)$$

## Expectation of the r.v $R_X$

To compute  $E[R_X]$ , we have:

$$E[R_X] = \sum_{i=1}^{n_1} E[R_i]$$

$$= n_1 \frac{N+1}{2}$$

$$= n_1 \frac{n_1 + n_2 + 1}{2}$$

### Variance of the r.v $R_X$

To compute  $var(R_X)$ , we have:

$$var(R_X) = \sum_{i=1}^{n_1} var(R_i) + \sum_{i=1}^{n_1} \sum_{i=1}^{n_1 - 1} cov(R_i, R_j)$$

where  $var(R_i) = \frac{N^2-1}{12}$  and  $cov(R_i, R_j) = -\frac{(N+1)}{12}$ . By replacing these results in the previous equation, we get

$$var(R_X) = n_1 \frac{N^2 - 1}{12} + n_1 (n_1 - 1) \left( -\frac{N+1}{12} \right)$$

$$= \frac{n_1}{12} \left[ N^2 - 1 - (n_1 - 1) (N+1) \right]$$

$$= \frac{n_1 (N+1)}{12} \left[ N - 1 - (n_1 - 1) \right]$$

$$= \frac{n_1 n_2 (N+1)}{12}$$

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### Distribution of the r.v $R_X$

So we have that

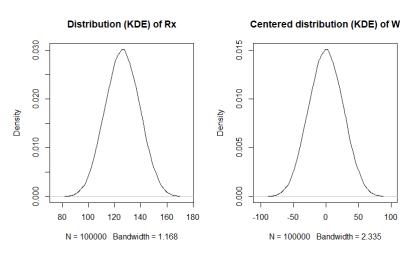
$$R_X \sim N\left(n_1 \frac{N+1}{2}, \frac{n_1 \ n_2 \ (N+1)}{12}\right)$$

This distribution is has obviously the same shape as the distribution of W since they both differ by the constant  $n_1(N+1)$ . See next slides.

## Empirical distribution of $R_X$ and W in R

```
1 # distribution of Rx and W under HO using simulation
 2 set. seed (2023)
 3 n = 100000
 4 X = matrix(rep(0, 12*n), nrow = n, ncol = 12)
 5 Y = matrix(rep(0, 8*n), nrow = n, ncol = 8)
 6 S = R = matrix(rep(0, (12+8)*n), nrow = n, ncol = 12+8)
 7 Rx = W = numeric(n)
8
9 for(i in 1:n) {
10 X[i,] = runif(12, min = 0, max = 2)
11 Y[i,] = runif(8, min = 0, max = 2)
12 S[i] = c(X[i], Y[i])
13 R[i] = rank(S[i])
14 Rx[i] = sum(R[i, 1:12])
15 W[i] = 2*Rx[i] + (12*(N+1))
16 }
17
18 # mean and variance of Rx and W
19 mean(Rx): var(Rx)
20 # [1] 126.048 about 12*(20+1)/2
21 # [1] 168.2862 about 12*8*(20+1)/12
22 mean(W): var(W)
23 # [1] 504.096 about (2*(12*(20+1)/2)) + 12*(20+1)
24 # [1] 673.145
25
26 # plots
27 \text{ par}(\text{mfrow} = c(1,2))
28 plot(density(Rx), main = 'Distribution (KDE) of Rx')
29 \text{ plot}(\text{density}(W - \text{rep}((2*(12*(20+1)/2)) + 12*(20+1), n)), main = 'Centered')
        distribution (KDE) of W')
```

## Density plots of $R_X$ and W



#### References

Bagdonavičius V., Kruopis J., Nikulin M. S., Non-parametric Tests for Complete Data (2011), Wiley, ISBN 978-1-84821-269-5 (hardback)

The R Project for Statistical Computing: https://www.r-project.org/

course notes