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#Question 2

#Rule to go by: If p is low, Reject H0. = null hypothesis

#Two independent samples

tA <- c(-0.15, 8.6, 5, 3.71, 4.29, 7.74, 2.48, 3.25, -1.15, 8.38)

tB <- c(2.55, 12.07, 0.46, 0.35, 2.69, -0.94, 1.73, 0.73, -0.35, -0.37)

par(mfrow = c(2,1))

hist(tA)

hist(tB)

par(mfrow = c(1,1))

median(tA) #4

median(tB) #0.595

#Wilcoxon-Mann Whitney Test

#####WILCOXON-MANN-WHITNEY TEST (TWO-SIDED)

?wilcox.test

wilcox.test(tA,tB, alternative = "two.sided", conf.int = TRUE, conf.level = 0.95) #p = 0.08921 Concludes the medians of t1 and t2 differ.

#The true location shift is not equal to 0, Both samples are not of similar shapes.

#Wilcoxon-Mann Whitney Test is inappropriate

#Siegel Tukey Test

library(DescTools)

?SiegelTukeyTest

SiegelTukeyTest(tA,tB, conf.int = TRUE, conf.level = 0.95)

#Siegel Tukey test is best used when medians are the same. That is not the case with these samples.

#This means the Siegel Tukey test, while more appropriate than Wilcoxon-Mann Whitney test, is not as appropriate as the Kolmogorov-Smirnov test.

lol yes, good job

show 1/2

#KS doesn't make any assumptions regarding the shape of the underlying distributions, so is the more appropriate test here. ✓

ks.test(x = tA, y = tB, alternative = "two.sided") #p = 0.05245

$H_0 + H_a ?$
Conclusion?

cdfA <- ecdf(tA)

cdfB <- ecdf(tB)

minMax <- seq(min(tA, tB), max(tA, tB), length.out=length(tA))

x0 <- minMax[which(abs(cdfA(minMax) - cdfB(minMax)) == max(abs(cdfA(minMax) - cdfB(minMax))))]

y0 <- cdfA(x0)

y1 <- cdfB(x0)

y0

y1

plot(cdfA, verticals=TRUE, col.points="blue", col="blue", main="Liquid Beverage Containers ECDFs", ylab="CDF", xlab="t")

lines(ecdf(tB), verticals=TRUE, col.points="green", col="green")

points(c(x0, x0), c(y0, y1), pch=16, col="red")

segments(x0, y0, x0, y1, col="red", lty="dotted")

#Both samples are not a similar shape & have different medians. ✓

#Only slightly so, but we fail to reject the null hypothesis.

#This test is the most appropriate.

#Therefore, we conclude The Kolmogorov-Smirnov is the most appropriate test to do here. ✓

	Treatment	Response	F^*	G	Abs. Difference
1	2	-1.15	0.1	0	0.1
2	2	-0.94	0.1	0.1	0
3	2	-0.57	0.1	0.2	0.1
4	2	-0.35	0.1	0.3	0.2
5	1	-0.15	0.2	0.3	0.1
6	2	0.35	0.2	0.4	0.2
7	2	0.96	0.2	0.5	0.3
8	2	0.73	0.2	0.6	0.4
9	2	1.73	0.2	0.7	0.5
10	1	2.48	0.3	0.7	0.4
11	2	2.55	0.3	0.8	0.5
12	2	2.69	0.3	0.9	0.6
13	1	3.25	0.4	0.9	0.5
14	1	3.71	0.5	0.9	0.4
15	1	4.24	0.6	0.9	0.3
16	1	5.00	0.7	0.9	0.2
17	1	7.74	0.8	0.9	0.1
18	1	8.38	0.9	0.9	0
19	1	8.60	1	0.9	0.1
20	2	12.07	1	1	0

$2.69 = b$, $0 = 0.6$ ✓ $D = \max_t |F(t) - G(t)| = |0.3 - 0.9| = 0.6$
 $G = 2.69$ $| -0.6 | = 0.6$

If we use R , we can confirm this

$n = 20$ $n_1 = 10$, $n_2 = 10$ two sided = 5 = answer

H_0 : We fail to reject H_0 & conclude that we have insufficient evidence of a difference between the CDFs of the ~~Responses~~ Responses/Samples

Critical values = 66, 70, 74, 80, 84, 87, 92, 94, 110, 110

#Q3

`tN <- c(74,79,92,65,253)`

`tS <- c(159,72,248,129,120)`

`median(tN)`

`median(tS)`

`hist(tN)`

`hist(tS)`

#Wilcoxon test is not appropriate

#Different shape & Median

#Siegel-Tukey test not appropriate

#Medians are not the same

#KS test is best, different shapes and medians are also different.

#KS doesn't make any assumptions regarding the shape of the underlying distributions,

#so is the more appropriate test here.

```
ks.test(Number, Symbol, alternative = "two.sided") #p-value = 0.3571
```

```
cdfN <- ecdf(tN)
```

```
cdfS <- ecdf(tS)
```

```
minMax <- seq(min(tN, tS), max(tN, tS), length.out=length(tN))
```

```
x0 <- minMax[which(abs(cdfN(minMax) - cdfS(minMax)) == max(abs(cdfN(minMax) -  
cdfS(minMax))))]
```

```
y0 <- cdfN(x0)
```

```
y1 <- cdfS(x0)
```

```
plot(cdfN, verticals=TRUE, col.points="blue", col="blue", main="Liquid Beverage Containers ECDFs",  
ylab="CDF", xlab="t")
```

```
lines(ecdf(tS), verticals=TRUE, col.points="green", col="green")
```

```
points(c(x0, x0), c(y0, y1), pch=16, col="red")
```

```
segments(x0, y0, x0, y1, col="red", lty="dotted")
```