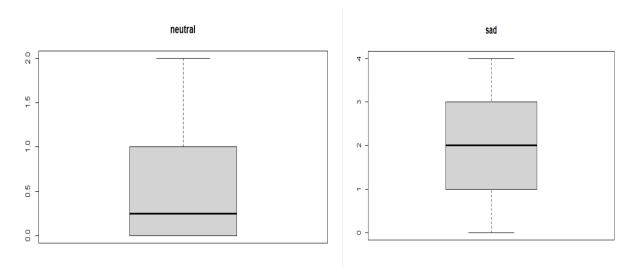
Harris Spahic 3/26/2021

"I pledge my honor I have abided by the Stevens Honor System"

Problem 7.71

A



The boxplots don't seem to be apx. normal but they don't seem
to have any outliers --> we can use t procedures

#B

```
> ntl_ss
[1] 14
> sad_ss
[1] 17
>
> ntl_mean
[1] 0.5714286
> sad_mean
[1] 2.117647
>
> ntl_sd
[1] 0.7300459
> sad_sd
[1] 1.244104
> | # H0 => mean(neural) = mean(sad)
#C # Ha => mean(neutral) != mean(sad)
```

```
Welch Two Sample t-test
 data: neutral and sad
 t = -4.3031, df = 26.48, p-value = 0.0002046
 alternative hypothesis: true difference in means is not equal to 0
 95 percent confidence interval:
  -2.2841749 -0.8082621
 sample estimates:
 mean of x mean of y
 0.5714286 2.1176471
#E
> t <- (ntl_mean - sad_mean) / sqrt(ntl_sd^2 / (ntl_ss - 1) + sad_sd^2 / (sad_ss - 1))
> SE <- sqrt(ntl_sd ^ 2 / ntl_ss + sad_sd ^ 2 / sad_ss)
> temp_Sp <- SE * 2.052
> interval <- c(ntl_mean - sad_mean - temp_Sp, ntl_mean - sad_mean + temp_Sp)</pre>
> interval
[1] -2.2835572 -0.8088798
> df <- df_t(ntl_sd, sad_sd, ntl_ss, sad_ss)</pre>
[1] 27
Problem 7.89 | #A
```

```
\# H0 \Rightarrow mean(bt) = mean(tmla)
# Ha => mean(bf) > mean(fmla)
> bf_ss
[1] 23
> fmla_ss
[1] 19
> bf_mean
[1] 13.3
> fmla_mean
[1] 12.4
> bf_sd
[1] 1.7
> fmla_sd
[1] 1.8
> df <- df_t(bf_sd, fmla_sd, bf_ss, fmla_ss)</pre>
> df
[1] 38
> t <- (bf_mean - fmla_mean) / sqrt(bf_sd \land 2 / bf_ss + fmla_sd \land 2 / fmla_ss)
[1] 1.653734
```

```
#B
SE <- sqrt(bf_sd \land 2 / bf_ss + fmla_sd \land 2 / fmla_ss)
1] 0.5442228
CI <- c(bf_mean - fmla_mean - SE * 2.042, bf_mean - fmla_mean + SE * 2.042)
1] -0.211303 2.011303
#C
We need both samples to be independent SRS from an Normal Distr.
Problem 7.102
#A
\cdot t <- (s1 - s2) / sqrt(s1 \wedge 2 / n1 + s2 \wedge 2 / n2)
1] -1.216636
pt(t, df) * 2
1] 0.2351046
> df <- df_t(s1, s2, n1, n2)
> df
[1] 25
```

#B

==> Value from table = 2.060

#C

```
# Since |-1.216626| < 2.060 --> Fail to reject HO
```

Problem 7.122 | #A

```
g1_ss
1] 10
g2_ss
1] 10
g1_mean
1] 49.692
g2_mean
1] 50.545
g1_sd
1] 2.317896
g2_sd
1] 1.92436
df <- df_t(g1_sd, g2_sd, g1_ss, g2_ss)
df
1] 18
 t \leftarrow (g1_{mean} - g2_{mean}) / sqrt(g1_{sd} \land 2 / g1_{ss} + g2_{sd} \land 2 / g2_{ss})
17 -0.8953783
pt(t, df) * 2
1] 0.3824027
#B
> dif_var
[1] 1.610668
> dif_mean
[1] -0.853
> df <- length(dif)
> df
[1] 10
> t \leftarrow (sum(dif) / df) / sqrt((sum(dif_sq) - (sum(dif) ^ 2) / df) / ((df - 1) * df))
[1] -2.125426
> df <- length(dif) - 1
> df
[1] 9
> pt(t, df)*2
[1] 0.06248424
```

P-value significantly smaller : 0.0624 | paired t-test < 0.3824 | 2 sample t-test Meaning very high chance to come to incorrect conclusion based on result

Problem 8.71 | #A

```
> f_prop
[1] 0.8
> m_prop
[1] 0.3939394
> f_SE
[1] 0.05163978
> m_SE
[1] 0.04252906
#B
std \leftarrow sqrt(f_prop*(1-f_prop) / fn + m_prop*(1-m_prop) / mn)
CI <- c(f_prop - m_prop - std * 1.645, f_prop - m_prop + std * 1.645)
CI
1] 0.2960128 0.5161084
#C
z <- (f_prop - m_prop) / SE
L] 5.220477
\#z = 5.2204 \gg 1.645 = alpha = 0.10 we reject the null hypothesis. Thus we
#have significant evidence to conclude the proportions are not equal.
```