Midterm1

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```
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr
           1.1.4
                       v readr
                                   2.1.5
## v forcats 1.0.0
                       v stringr 1.5.1
## v ggplot2 3.5.1
                      v tibble
                                   3.2.1
## v lubridate 1.9.4
                                   1.3.1
                       v tidyr
              1.0.4
## v purrr
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(car)
## Loading required package: carData
## Attaching package: 'car'
## The following object is masked from 'package:dplyr':
##
##
      recode
## The following object is masked from 'package:purrr':
##
##
       some
library(MVN)
library(Hotelling)
## Loading required package: corpcor
## Attaching package: 'Hotelling'
## The following object is masked from 'package:dplyr':
##
##
      summarise
```

```
set.seed(123)
load("Data/plasticDat.RData")
dat <- as_tibble(dat)</pre>
head(dat)
## # A tibble: 6 x 6
## setting
            V1
                  V2 tearResist gloss opacity
##
      <dbl> <int> <int> <dbl> <dbl> <dbl>
## 1
                         6.5 9.5
                                       4.4
       0 0 0
## 2
       0
             0 0
                         6.2 9.9
                                       6.4
       0 0 0 0
## 3
                         5.8 9.6
                                       3
                         6.5 9.6
## 4
                                       4.1
## 5
       0 0 0
                         6.5 9.2
                                     0.8
## 6
       1 0 1
                         6.9 9.1
                                       5.7
\mathbf{a}
load("Data/plasticDat.RData")
names(dat)
## [1] "setting"
                 "V1"
                            "V2"
                                       "tearResist" "gloss"
## [6] "opacity"
str(dat)
## 'data.frame': 20 obs. of 6 variables:
## $ setting : num 0 0 0 0 0 1 1 1 1 1 ...
## $ V1
            : int 0000000000...
## $ V2
          : int 0000011111...
## $ tearResist: num 6.5 6.2 5.8 6.5 6.5 6.9 7.2 6.9 6.1 6.3 ...
## $ gloss : num 9.5 9.9 9.6 9.6 9.2 9.1 10 9.9 9.5 9.4 ...
## $ opacity : num 4.4 6.4 3 4.1 0.8 5.7 2 3.9 1.9 5.7 ...
head(dat)
    setting V1 V2 tearResist gloss opacity
## 1
        0 0 0
                     6.5 9.5
                                  4.4
         0 0 0
## 2
                      6.2 9.9
                                  6.4
## 3
        0 0 0
                    5.8 9.6
                                  3.0
## 4
        0 0 0
                    6.5 9.6
                                 4.1
     0 0 0 6.5 9.2
1 0 1 6.9 9.1
## 5
                                  0.8
## 6
                                  5.7
```

```
man <- manova(cbind(tearResist, gloss, opacity) ~ setting, data=dat)</pre>
man_sum <- summary(man, test="Wilks")</pre>
print(man_sum)
##
                  Wilks approx F num Df den Df
              1 0.34858
                         9.9666
                                      3
                                            16 0.0006044 ***
## setting
## Residuals 18
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
pval <- man_sum$stats["setting","Pr(>F)"]
cat("MANOVA p-value =", pval, "\n")
## MANOVA p-value = 0.0006044414
b
library(biotools)
## Loading required package: MASS
##
## Attaching package: 'MASS'
## The following object is masked from 'package:dplyr':
##
##
       select
## ---
## biotools version 4.3
print(boxM(dat[,c("tearResist","gloss","opacity")], dat$setting))
##
## Box's M-test for Homogeneity of Covariance Matrices
## data: dat[, c("tearResist", "gloss", "opacity")]
## Chi-Sq (approx.) = 24.015, df = 18, p-value = 0.1546
res <- residuals(man)
sw <- sapply(as.data.frame(res), function(x) shapiro.test(x)$p.value)</pre>
print(sw)
## tearResist
                   gloss
                            opacity
## 0.2913991 0.5008235 0.4656299
```

c, d

```
settings <- sort(unique(dat$setting))</pre>
pairs
       <- combn(settings, 2, simplify = FALSE)</pre>
raw_p <- sapply(pairs, function(p) {</pre>
 mv <- manova(cbind(tearResist, gloss, opacity) ~ factor(setting),</pre>
                data = subset(dat, setting %in% p))
  sum <- summary(mv, test = "Wilks")</pre>
  sum$stats[1, "Pr(>F)"]
})
adj_p <- p.adjust(raw_p, method = "bonferroni")</pre>
pairwise_df <- data.frame(</pre>
  Comparison = sapply(pairs, function(p) paste(p, collapse = " vs ")),
  raw_p_value = raw_p,
  adj_p_value = adj_p
pairwise_df
     Comparison raw_p_value adj_p_value
## 1
         0 vs 1 0.636585195 1.00000000
## 2
         0 vs 2 0.067731972 0.40639183
        0 vs 3 0.005562226 0.03337336
## 3
        1 vs 2 0.093991698 0.56395019
       1 vs 3 0.118179705 0.70907823
## 5
## 6
         2 vs 3 0.014552976 0.08731785
\mathbf{e}
sub03 <- subset(dat, setting %in% c("0","3"))</pre>
lapply(sub03[c("tearResist","gloss","opacity")], function(x) {
  t.test(x ~ setting, data = sub03)
})
## $tearResist
##
## Welch Two Sample t-test
## data: x by setting
## t = -5.4444, df = 7.7679, p-value = 0.0006773
## alternative hypothesis: true difference in means between group 0 and group 3 is not equal to 0
## 95 percent confidence interval:
## -1.3972491 -0.5627509
## sample estimates:
## mean in group 0 mean in group 3
##
              6.30
                               7.28
```

```
##
##
## $gloss
##
##
  Welch Two Sample t-test
##
## data: x by setting
## t = 0.63444, df = 5.8628, p-value = 0.5497
## alternative hypothesis: true difference in means between group 0 and group 3 is not equal to 0
## 95 percent confidence interval:
## -0.4606059 0.7806059
## sample estimates:
## mean in group 0 mean in group 3
##
              9.56
                              9.40
##
##
## $opacity
##
  Welch Two Sample t-test
##
## data: x by setting
## t = -0.83551, df = 7.4075, p-value = 0.4296
## alternative hypothesis: true difference in means between group 0 and group 3 is not equal to 0
## 95 percent confidence interval:
## -4.86258 2.30258
## sample estimates:
## mean in group 0 mean in group 3
              3.74
                              5.02
print(summary.aov(man))
## Response tearResist :
##
              Df Sum Sq Mean Sq F value
                                            Pr(>F)
## setting
               1 2.4649 2.46490 24.641 0.0001004 ***
## Residuals
             18 1.8006 0.10003
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Response gloss :
##
              Df Sum Sq Mean Sq F value Pr(>F)
## setting
               1 0.4489 0.44890 1.7427 0.2033
## Residuals
              18 4.6366 0.25759
## Response opacity:
##
              Df Sum Sq Mean Sq F value Pr(>F)
## setting
               1 2.465 2.4649 0.6185 0.4419
## Residuals
             18 71.741 3.9856
tw <- manova(cbind(tearResist, gloss, opacity) ~ V1 * V2, data=dat)</pre>
print(summary(tw, test="Wilks"))
                  Wilks approx F num Df den Df
## V1
              1 0.38186 7.5543
                                           14 0.003034 **
                                      3
```