

HW 3

Q1.1

```
data = read.delim("multiple.txt", sep = " ", header=FALSE)
```

```
sample_mean = colMeans(data)
sample_sd = sapply(data, sd)
hnull = hnull = integer(50)
```

```
n = nrow(data)
t_statistic = (sample_mean - hnull)/(sample_sd/sqrt(n))
```

```
pval = 2*apply(rbind(pt(t_statistic, df = n-1, lower.tail = F),
                      pt(t_statistic, df = n-1, lower.tail = T)), 2, min)
rejection_p <- pval < 0.1
print(pval)
```

```
##          V1          V2          V3          V4          V5          V6
## 7.477468e-34 1.030644e-32 1.613953e-33 5.107798e-37 4.299681e-35 4.181053e-33
##          V7          V8          V9          V10          V11          V12
## 1.782424e-39 2.281744e-34 4.767758e-34 9.020768e-42 8.501800e-01 3.387655e-01
##          V13          V14          V15          V16          V17          V18
## 8.433591e-01 2.353081e-01 8.346443e-01 3.410084e-01 4.495285e-01 8.988573e-01
##          V19          V20          V21          V22          V23          V24
## 4.855360e-01 8.298508e-02 7.435778e-01 4.344605e-01 5.521855e-01 4.775194e-01
##          V25          V26          V27          V28          V29          V30
## 7.573900e-01 1.354197e-01 9.927262e-01 2.944327e-01 6.783256e-01 1.898939e-01
##          V31          V32          V33          V34          V35          V36
## 4.716828e-01 9.302681e-02 4.736660e-01 9.189993e-01 2.138515e-01 6.519921e-01
##          V37          V38          V39          V40          V41          V42
## 3.408716e-01 2.211123e-01 8.806238e-01 7.315268e-02 7.171567e-01 6.472996e-01
##          V43          V44          V45          V46          V47          V48
## 6.915055e-01 2.475067e-02 2.191321e-01 7.678119e-01 3.489369e-01 1.367531e-01
##          V49          V50
## 8.886784e-01 7.316609e-01
```

```
print(rejection_p)
```

```
##  V1  V2  V3  V4  V5  V6  V7  V8  V9  V10  V11  V12  V13
## TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE FALSE FALSE FALSE
## V14 V15 V16 V17 V18 V19 V20 V21 V22 V23 V24 V25 V26
## FALSE FALSE FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE FALSE FALSE FALSE
## V27 V28 V29 V30 V31 V32 V33 V34 V35 V36 V37 V38 V39
## FALSE FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## V40 V41 V42 V43 V44 V45 V46 V47 V48 V49 V50
## TRUE FALSE FALSE FALSE TRUE FALSE FALSE FALSE FALSE FALSE FALSE
```

Q1.2

```
alpha = 0.1
FWER = 1 - (1 - alpha)^50
R = sum(rejection_p)
V = sum(rejection_p[11:50])
print(FWER)
```

```
## [1] 0.9948462
```

```
print(V/R)
```

```
## [1] 0.2857143
```

FWER: 0.9948462 FDP: 0.2857143

Q1.3

```
pval.bon = p.adjust(pval, method = "bonferroni")
rejection_bon = pval.bon < 0.1
print(rejection_bon)
```

```
##      V1      V2      V3      V4      V5      V6      V7      V8      V9      V10     V11     V12     V13
## TRUE  TRUE  TRUE  TRUE  TRUE  TRUE  TRUE  TRUE  TRUE  TRUE FALSE FALSE FALSE
##      V14     V15     V16     V17     V18     V19     V20     V21     V22     V23     V24     V25     V26
## FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
##      V27     V28     V29     V30     V31     V32     V33     V34     V35     V36     V37     V38     V39
## FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
##      V40     V41     V42     V43     V44     V45     V46     V47     V48     V49     V50
## FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

```
FWER.bon = 1 - (1 - alpha/50)^50
print(FWER.bon)
```

```
## [1] 0.09525318
```

FWER: 0.09525318

Q1.4

```
pval.bh = p.adjust(pval, method = "BH")
rejection_bh = pval.bh < 0.1
alphas = 0.1 * seq(1/50, 1, by = 1/50)
FWER.bh = 1 - prod(1 - alphas)
FDP_bh = sum(rejection_bh[11:50])/sum(rejection_bh)
print(FWER.bh)
```

```
## [1] 0.928672
```

```
print(FDP_bh)
```

```
## [1] 0
```

BH's FWER is slightly lower than step1 and way higher than step2 but BH's FDP is 0, which is way lower than step1 and 3.

Q2

The null hypothesis is that the population means of all six variables are the same for the genuine and counterfeit bank notes. The alternative hypothesis is that they are different. We can assume that they have the same population variance-covariance matrices.

```
library(glue)
banknotes <- read.delim("SwissBankNotes.txt", sep = "")
df <- banknotes
genuine <- df[0:100,]
counterfeit <- df[101:200,]
n <- 100
delta <- colMeans(genuine) - colMeans(counterfeit)
p <- 6
Sg <- cov(genuine)
Sc <- cov(counterfeit)
S_pooled <- ((n-1)*Sg + (n-1)*Sc) / (n+n-2)
t_squared <- (n*n)/(n+n) * t(delta) %*% solve(S_pooled) %*% (delta)
f_statistic <- t_squared * (n+n-p-1)/(p*(n+n-2))
cv <- qf(0.05, p, n+n-p-1, lower.tail = FALSE)
print(glue("The F statistic is {f_statistic} and the critical value is {cv}"))
```

```
## The F statistic is 391.921702277771 and the critical value is 2.14580146767029
```

Since the F statistic is larger than the critical value, we are able to reject the null hypothesis. We have evidence that the counterfeits are distinguishable from the genuine bank notes on at least one of the variables.