HW2

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Chapter 4

10

```
#a
mu < -c(3,1,4)
SIGMA \leftarrow-rbind(c(6,1,-2),c(1,13,4),c(-2,4,4))
c < -c(2,-1,3)
(zmu<-c%*%mu)
## [,1]
## [1,] 17
(zSIGMA<-t(c)%*%SIGMA%*%c)
## [,1]
## [1,] 21
# z~(17,21)
cb < -rbind(c(1,1,1),c(1,-1,2))
(zbmu<-cb%*%mu)
##
      [,1]
## [1,] 8
## [2,]
        10
(zbSIGMA<-cb%*%SIGMA%*%t(cb))
##
      [,1] [,2]
## [1,] 29 -1
## [2,] -1
# z1,z2~N2((8,10),[(29,-1),(-1,9)])
# y2~N(1,13)
\#d
# y1,y3~N2((3,4),[(6,-2),(-2,4)])
#e
C \leftarrow rbind(c(1,0,0),c(0,0,1),c(.5,.5,0))
(cmu<-C%*%mu)
##
      [,1]
## [1,] 3
## [2,]
```

```
## [3,]
(cSIGMA<-C%*%SIGMA%*%t(C))
     [,1] [,2] [,3]
## [1,] 6.0 -2 3.50
## [2,] -2.0
             4 1.00
## [3,] 3.5
             1 5.25
# y1,y3,.5(y1+y2)~N3((3,4,2),[(6,-2,3.5),(-2,4,1),(3.5,1,5.25)])
```

11

```
#b
SIGMARoot <- eigen(SIGMA)$vectors%*%diag(sqrt(eigen(SIGMA)$values))%*%t(eigen(SIGMA)$vectors)
(SIGMARootInverse<-solve(SIGMARoot))
##
               [,1]
                           [,2]
                                      [,3]
## [1,] 0.46496849 -0.06966936 0.1701484
## [2,] -0.06966936  0.32645938 -0.1657086
## [3,] 0.17014841 -0.16570861 0.6916013
# z=SIGMARootInverse\%*\%c(y-3,y-1,y-4)
# Distributed chi-square(3) because statistical distance is distributed chi-square(p)
```

14

```
#a
mu < -c(2, -3, 4)
(SIGMA \leftarrow rbind(c(4,-3,0),c(-3,6,0),c(0,0,5)))
        [,1] [,2] [,3]
## [1,]
             -3
        4
## [2,]
         -3
             6
                     0
## [3,]
          0
               0
                     5
# Dependent due to covariance not equal to O
#b
# Independent due to covariance equal to 0
# Independent due to covariance equal to 0
c < -rbind(c(1,1,0),c(0,0,1))
(dSIGMA<-c%*%SIGMA%*%t(c))
##
      [,1] [,2]
## [1,] 4 0
        0 5
## [2,]
```

```
# Independent due to covariance equal to 0

#e

c<-rbind(c(1,0,1),c(0,1,0))

(eSIGMA<-c%*%SIGMA%*%t(c))

## [,1] [,2]

## [1,] 9 -3

## [2,] -3 6

# Dependent due to covariance not equal to 0
```

16

```
#a
mu < -c(2,-1,3,1)
SIGMA \leftarrow rbind(c(7,3,-3,2),c(3,6,0,4),c(-3,0,5,-2),c(2,4,-2,4))
(n<-mu[1:2])
## [1] 2 -1
(z<-SIGMA[1:2,3:4]%*%solve(SIGMA[3:4,3:4]))
        [,1] [,2]
## [1,] -0.5 0.25
## [2,] 0.5 1.25
\# n+z\%*\%c(x1-3,x2-1)
(SIGMA[1:2,1:2]-SIGMA[1:2,3:4]%*%solve(SIGMA[3:4,3:4])%*%SIGMA[3:4,1:2])
##
        [,1] [,2]
## [1,]
          5
## [2,]
           2
                1
```

Chapter 5

11

```
nullvec<-c(6,11)
Y<-cbind(c(3,6,5,10),c(10,12,14,9))
meanvec<-colMeans(Y)
varmat<-var(Y)
n=nrow(Y)
p=ncol(Y)
(T2 <- n*t(meanvec-nullvec)%*%solve(varmat)%*%(meanvec-nullvec))

## [1,]
## [1,] 0.06103286
(critval <- (n-1)*p/(n-p)*qf(.95,p,n-p))
## [1] 57</pre>
```

```
(T2>critval)
## [,1]
## [1,] FALSE
# Fail to reject HO

12
probe <- read.table("T3_6_PROBE.dat")</pre>
```

```
probe<-probe[,-1]</pre>
#a
nullvec < -c(30, 25, 40, 25, 30)
meanvec <- colMeans(probe)</pre>
varmat <- var(probe)</pre>
n=nrow(probe)
p=ncol(probe)
(T2 <- n*t(meanvec-nullvec)%*%solve(varmat)%*%(meanvec-nullvec))
           [,1]
## [1,] 85.3327
(critval <- (n-1)*p/(n-p)*qf(.95,p,n-p))
## [1] 36.56145
(T2>critval)
##
        [,1]
## [1,] TRUE
# reject HO
#b
t.test(probe$V2,mu=nullvec[1])
##
## One Sample t-test
##
## data: probe$V2
## t = 2.5039, df = 10, p-value = 0.03124
## alternative hypothesis: true mean is not equal to 30
## 95 percent confidence interval:
## 30.67082 41.51099
## sample estimates:
## mean of x
## 36.09091
# reject HO, true mean not equal to 30
t.test(probe$V3,mu=nullvec[2])
##
##
   One Sample t-test
```

```
## One Sample t-test
##
## data: probe$V3
## t = 0.26652, df = 10, p-value = 0.7953
```

```
## alternative hypothesis: true mean is not equal to 25
## 95 percent confidence interval:
## 20.98542 30.10549
## sample estimates:
## mean of x
## 25.54545
# Fail to reject HO, true mean equal to 25
t.test(probe$V4,mu=nullvec[3])
##
##
   One Sample t-test
##
## data: probe$V4
## t = -2.5157, df = 10, p-value = 0.03061
## alternative hypothesis: true mean is not equal to 40
## 95 percent confidence interval:
## 28.85722 39.32460
## sample estimates:
## mean of x
## 34.09091
# reject HO, true mean not equal to 40
t.test(probe$V5,mu=nullvec[4])
##
##
   One Sample t-test
##
## data: probe$V5
## t = 0.95104, df = 10, p-value = 0.364
## alternative hypothesis: true mean is not equal to 25
## 95 percent confidence interval:
## 21.94811 32.59735
## sample estimates:
## mean of x
## 27.27273
# Fail to reject HO, true mean equal to 25
t.test(probe$V6,mu=nullvec[5])
##
##
   One Sample t-test
##
## data: probe$V6
## t = 0.31613, df = 10, p-value = 0.7584
## alternative hypothesis: true mean is not equal to 30
## 95 percent confidence interval:
## 25.60131 35.85323
## sample estimates:
## mean of x
## 30.72727
# Fail to reject HO, true mean equal to 30
```

```
fBeetles <- read.table("T5_5_FBEETLES.dat")
fBeetles<-fBeetles[,-1]
#a
g1<- subset(fBeetles, V2==1)[,-1]
g2<- subset(fBeetles, V2==2)[,-1]
p <- ncol(g1)
n1 \leftarrow nrow(g1)
n2 \leftarrow nrow(g2)
mean1 <- colMeans(g1)</pre>
mean2 <- colMeans(g2)</pre>
S1 \leftarrow var(g1)
S2 \leftarrow var(g2)
Sg \leftarrow ((n1-1)*S1+(n2-1)*S2)/(n1+n2-2)
(T2 <- n1*n2/(n1+n2)*t(mean1-mean2)%*%solve(Sg)%*%(mean1-mean2))
##
             [,1]
## [1,] 133.4873
a \leftarrow p*(n1+n2-2)/(n1+n2-p-1)
(crit.val <- a*qf(.95,p,n1+n2-p-1))
## [1] 11.53483
T2>crit.val
         [,1]
## [1,] TRUE
# Reject HO
#b
p.vals \leftarrow rep(1,p)
t.stat <- (mean1[1]-mean2[1])/sqrt((1/n1+1/n2)*Sg[1,1])
p.vals[1] <- 2*pt(abs(t.stat), n1+n2-2, lower.tail = F)</pre>
t.stat \leftarrow (mean1[2]-mean2[2])/sqrt((1/n1+1/n2)*Sg[2,2])
p.vals[2] \leftarrow 2*pt(abs(t.stat), n1+n2-2, lower.tail = F)
t.stat <- (mean1[3]-mean2[3])/sqrt((1/n1+1/n2)*Sg[3,3])
p.vals[3] \leftarrow 2*pt(abs(t.stat), n1+n2-2, lower.tail = F)
t.stat <- (mean1[4]-mean2[4])/sqrt((1/n1+1/n2)*Sg[4,4])
p.vals[4] \leftarrow 2*pt(abs(t.stat), n1+n2-2, lower.tail = F)
(p.vals)
## [1] 4.049179e-04 4.326232e-04 1.645393e-06 1.236427e-05
# Reject HO for all
solve(Sg)%*%(mean1-mean2)
             [,1]
## V3 0.3452490
## V4 -0.1303878
## V5 -0.1064338
## V6 -0.1433533
```

```
goods <- read.table("T5_8_GOODS.dat")</pre>
goods<-goods[,-1]
g1<- subset(goods, V2==1)[,-1]
g2<- subset(goods, V2==2)[,-1]
p <- ncol(g1)
n1 <- nrow(g1)
n2 <- nrow(g2)
mean1 <- colMeans(g1)</pre>
mean2 <- colMeans(g2)</pre>
S1 \leftarrow var(g1)
S2 \leftarrow var(g2)
Sg \leftarrow ((n1-1)*S1+(n2-1)*S2)/(n1+n2-2)
(T2 \leftarrow n1*n2/(n1+n2)*t(mean1-mean2)%*%solve(Sg)%*%(mean1-mean2))
              [,1]
## [1,] 18.46248
a \leftarrow p*(n1+n2-2)/(n1+n2-p-1)
(crit.val <- a*qf(.95,p,n1+n2-p-1))
## [1] 15.11664
T2>crit.val
##
         [,1]
## [1,] TRUE
# Reject HO
solve(Sg)%*%(mean1-mean2)
                [,1]
## V3 -0.056896013
## V4 -0.009709542
## V5 -0.242134127
## V6 -0.071282739
21
essay <- read.table("T5_9_ESSAY.dat")</pre>
essay <- essay[,-1]
#a
dword<-as.matrix(essay[,1]-essay[,3])</pre>
dverb<-as.matrix(essay[,2]-essay[,4])</pre>
diff <-cbind(dword,dverb)</pre>
p<-ncol(diff)</pre>
n<-nrow(diff)</pre>
meandiff <- colMeans(diff)</pre>
varmat <- var(diff)</pre>
(T2 <- n*t(meandiff)%*%solve(varmat)%*%meandiff)</pre>
              [,1]
##
```

```
## [1,] 15.19123
(critval \leftarrow p*(n-1)/(n-p)*qf(.95,p,n-p))
## [1] 8.196602
T2>critval
##
        [,1]
## [1,] TRUE
#reject HO
#b
solve(varmat)%*%meandiff
##
               [,1]
## [1,] -0.03601558
## [2,] 0.04770629
t.test(dword)
##
## One Sample t-test
##
## data: dword
## t = -3.8371, df = 14, p-value = 0.001813
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## -51.13367 -14.46633
## sample estimates:
## mean of x
##
       -32.8
# Reject HO
t.test(dverb)
##
##
  One Sample t-test
##
## data: dverb
## t = -2.4362, df = 14, p-value = 0.0288
\#\# alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## -6.644008 -0.422659
## sample estimates:
## mean of x
## -3.533333
# Reject HO
```