

MVA HW#4

Problem 6.7

```
> S1 = matrix(c(13825.3,23823.4,23823.4,73107.4),2,2)
> S2 = matrix(c(8632.0,19616.7,19616.7,55964.5),2,2)
> n1 = 45
> n2 = 55
> p = 2
> xbar1 = c(204.4,556.6)
> xbar2 = c(130,355.0)
> TSquare.two_sample.withSummary(mean1=xbar1, mean2=xbar2,
cov1 = S1, cov2 = S2, n1=n1, n2=n2, p=p, pooled = TRUE)
-----
Method: Two Population T-square Test with Equal Variance (P
ooled)
-----
Pooled covariance:
      [,1] [,2]
[1,] 10963.69 21505.42
[2,] 21505.42 63661.31
-----
T-square value: 16.06622
P-value: 0.0006343777
```

Problem 6.8

```
> x1 = c(6,5,8,4,7,3,1,2,2,5,3,2)
> x2 = c(7,9,6,9,9,3,6,3,3,1,1,3)
> trt = c(rep(1,5),rep(2,3),rep(3,4))
> MANOVA(cbind(x1,x2), trt)
-----
MANOVA Procedure
-----
Grand Mean:
x1 x2
 4  5

Treatment Means:
      group 1 group 2 group 3
x1      6      2      3
x2      8      4      2

Treatment Effects:
      group 1 group 2 group 3
x1      2      -2      -1
x2      3      -1      -3

Residuals:
      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11]
1] [,12]
[1,]  0    -1    2    -2    1    1    -1    0    -1    2
     0    -1
```

Problem 6.21

```
> n1 = 20
> n2 = 20
> p = 4
> xbar1 = c(2.287, 12.6,0.347, 14.830)
> xbar2 = c(2.404,7.155,0.524,12.840)
> S1 = matrix(c(0.459,.254,-.026,-.244,.254,27.465,-.589,-.
267,-.026,-.589,.030,.102,-.244,-.267,.102,6.854),4,4)
> S2 = matrix(c(.944,-.089,.002,-.719,-.089,16.432,-.4,19.0
44,.002,-.400,.024,-.094,-.719,19.044,-.094,61.854),4,4)
```

```
-----
Confident Intervals at 95% confident level:
Simontaneous CI:
      low      high
1 21.80733 126.9927
2 74.86846 328.3315

Bonferroni CI:
      low      high
1 42.64506 106.1549
2 125.08073 278.1193
-----
The most critical linear combination to reject H0 is:
      [,1]
[1,] 0.00170252
[2,] 0.00259163
-----

[2,] -1    1    -2    1    1    -1    2    -1    1    -1
-1    1

-----
The MANOVA Table Elements:
-----
Sstrt (df = 2)
      x1 x2
[1,] 36 48
[2,] 48 84

SSerror (df = 9)
      [,1] [,2]
[1,] 18 -13
[2,] -13 18

SStotal (df = 11)
      x1 x2
[1,] 54 35
[2,] 35 102
-----
Wilk's Test:
-----
lambda = 0.03618959      ratio = 4.256639      p-value:
0.01554857
Bartlett's correction: 28.21136      p-value: 1.13011e-05

> TSquare.two_sample.withSummary(mean1=xbar1, mean2=xbar2,
cov1 = S1, cov2 = S2, n1=n1, n2=n2, p=p, pooled = TRUE)
-----
Method: Two Population T-square Test with Equal Variance (P
ooled)
-----
Pooled covariance:
      [,1] [,2] [,3] [,4]
[1,] 0.7015 0.0825 -0.0120 -0.4815
[2,] 0.0825 21.9485 -0.4945 9.3885
```

```
[3,] -0.0120 -0.4945 0.0270 0.0040
[4,] -0.4815 9.3885 0.0040 34.3540
```

T-square value: 15.82996
P-value: 0.01386451

Confident Intervals at 95% confident level:
Simontaneous CI:

	low	high
1	-1.0140653	0.780065294
2	0.4272082	10.462791829
3	-0.3529917	-0.001008335
4	-4.2876756	8.267675648

Bonferroni CI:

	low	high
1	-0.5458907	0.31189075
2	3.0459723	7.84402771
3	-0.2611424	-0.09285763
4	-1.0113835	4.99138355

The most critical linear combination to reject H0 is:

	[,1]
[1,]	-0.24176170
[2,]	0.16009566
[3,]	-3.73254518
[4,]	0.01122035

```
> TSquare.two_sample.withSummary(mean1=xbar1, mean2=xbar2,
cov1 = S1, cov2 = S2, n1=n1, n2=n2, p=p, pooled = FALSE)
```

Method: Two Population T-square Test with Unequal Variance
(Unpooled)

Problem 6.24

```
> x1 = c(131,125,131,119,136,138,139,125,131,134,124,133,13
8,148,126,135,132,133,131,133,132,133,138,130,136,134,136,1
33,138,138)
> x2 = c(138,131,132,132,143,137,130,136,134,134,138,134,13
4,129,124,136,145,130,134,125,130,131,137,127,133,123,137,1
31,133,133)
> x3 = c(89,92,99,96,100,89,108,93,102,99,101,97,98,104,95,
98,100,102,96,94,91,100,94,99,91,95,101,96,100,91)
> x4 = c(49,48,50,44,54,56,48,48,51,51,48,48,45,51,45,52,5
4,48,50,46,52,50,51,45,49,52,54,49,55,46)
> trt = c(rep(1,10),rep(2,10),rep(3,10))
> x = cbind(x1,x2,x3,x4)
> MANOVA(X,trt)
```

MANOVA Procedure

Grand Mean:

	x1	x2	x3	x4
133.00000	133.03333	97.00000	49.63333	

Treatment Means:

	group 1	group 2	group 3
x1	130.9	133.3	134.8
x2	134.7	132.9	131.5
x3	96.7	98.5	95.8
x4	49.9	48.7	50.3

Treatment Effects:

Unpooled covariance:

	[,1]	[,2]	[,3]	[,4]
[1,]	0.07015	0.00825	-0.00120	-0.04815
[2,]	0.00825	2.19485	-0.04945	0.93885
[3,]	-0.00120	-0.04945	0.00270	0.00040
[4,]	-0.04815	0.93885	0.00040	3.43540

T-square value: 15.82996
P-value: 0.003256029

Confident Intervals at 95% confident level:
Simontaneous CI:

	low	high
1	-0.9328212	0.69882118
2	0.8816523	10.00834768
3	-0.3370527	-0.01694729
4	-3.7191282	7.69912816

Bonferroni CI:

	low	high
1	-0.5355858	0.30158576
2	3.1036139	7.78638610
3	-0.2591207	-0.09487933
4	-0.9392691	4.91926909

The most critical linear combination to reject H0 is:

	[,1]
[1,]	-2.4176170
[2,]	1.6009566
[3,]	-37.3254518
[4,]	0.1122035

	group 1	group 2	group 3
x1	-2.1000000	0.3000000	1.8000000
x2	1.6666667	-0.1333333	-1.5333333
x3	-0.3000000	1.5000000	-1.2000000
x4	0.2666667	-0.9333333	0.6666667

Residuals:

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]	[,11]	[,12]	[,13]	[,14]	[,15]	[,16]	[,17]	[,18]	[,19]	[,20]	[,21]	[,22]
[1,]	0.1	-5.9	0.1	-11.9	5.1	7.1	8.1	-5.9	0.1	3.1	-9.3	-0.3	4.7	14.7	-7.3	1.7	-1.3	-0.3	-2.3	-0.3	-2.8	-1.8
[2,]	3.3	-3.7	-2.7	-2.7	8.3	2.3	-4.7	1.3	-0.7	-0.7	5.1	1.1	1.1	-3.9	-8.9	3.1	12.1	-2.9	1.1	-7.9	-1.5	-0.5
[3,]	-7.7	-4.7	2.3	-0.7	3.3	-7.7	11.3	-3.7	5.3	2.3	2.5	-1.5	-0.5	5.5	-3.5	-0.5	1.5	3.5	-2.5	-4.5	-4.8	4.2
[4,]	-0.9	-1.9	0.1	-5.9	4.1	6.1	-1.9	-1.9	1.1	1.1	-0.7	-0.7	-3.7	2.3	-3.7	3.3	5.3	-0.7	1.3	-2.7	1.7	-0.3
	[,23]	[,24]	[,25]	[,26]	[,27]	[,28]	[,29]	[,30]														
[1,]	3.2	-4.8	1.2	-0.8	1.2	-1.8	3.2	3.2														
[2,]	5.5	-4.5	1.5	-8.5	5.5	-0.5	1.5	1.5														
[3,]	-1.8	3.2	-4.8	-0.8	5.2	0.2	4.2	-4.8														
[4,]	0.7	-5.3	-1.3	1.7	3.7	-1.3	4.7	-4.3														

The MANOVA Table Elements:

ssrt (df = 2)

	x1	x2	x3	x4
[1,]	77.4	-63.000000	-10.8	3.600000
[2,]	-63.0	51.466667	11.4	-4.533333
[3,]	-10.8	11.400000	37.8	-22.800000
[4,]	3.6	-4.533333	-22.8	13.866667

sserror (df = 27)

	[,1]	[,2]	[,3]	[,4]
--	------	------	------	------

```
[1,] 822.6 90.0 186.8 220.4
[2,] 90.0 651.5 1.6 217.9
[3,] 186.8 1.6 570.2 56.8
[4,] 220.4 217.9 56.8 277.1
```

```
SStotal (df = 29)
```

```
      x1      x2      x3      x4
[1,] 900 27.0000 176 224.0000
[2,] 27 702.9667 13 213.3667
[3,] 176 13.0000 608 34.0000
[4,] 224 213.3667 34 290.9667
```

```
-----
```

```
Wilk's Test:
```

```
-----
```

```
lambda = 0.7020877      ratio = 0.1934503      p-value:
0.9407504
```

```
Bartlett's correction: 3.006424      p-value: 0.9339537
```

```
> cor(x[1:10,])
```

```
      x1      x2      x3      x4
x1 1.0000000 0.2890107 0.3419187 0.7498376
x2 0.2890107 1.0000000 -0.3077189 0.6098596
x3 0.3419187 -0.3077189 1.0000000 -0.1044035
x4 0.7498376 0.6098596 -0.1044035 1.0000000
```

```
> cor(x[11:20,])
```

```
      x1      x2      x3      x4
x1 1.0000000 -0.1211279 0.4459786 0.2627420
x2 -0.1211279 1.0000000 0.3302588 0.6757719
x3 0.4459786 0.3302588 1.0000000 0.4532676
x4 0.2627420 0.6757719 0.4532676 1.0000000
```

```
> cor(x[21:30,])
```

```
      x1      x2      x3      x4
x1 1.0000000 0.64895539 -0.16062260 0.3521922
x2 0.6489554 1.00000000 0.05200949 0.2493121
x3 -0.1606226 0.05200949 1.00000000 0.3160608
x4 0.3521922 0.24931209 0.31606076 1.0000000
```

Define functions:

1.

```
> TSquare.two_sample.withSummary = function(mean1, mean2, cov1, cov2, n1, n2, p, mu = 0, x1_minus_x2=TRUE, alpha = 0.05, pooled =
FALSE){
+   if(x1_minus_x2){
+     mean_diff = mean1 - mean2
+   }else{
+     mean_diff = mean1 + mean2
+   }
+   diff = mean_diff - mu
+   cat('-----\n')
+   if(pooled){
+     cat("Method: Two Population T-square Test with Equal Variance (Pooled)\n-----\n")
+     cov.pooled = ((n1-1)*cov1+(n2-1)*cov2)/(n1+n2-2)
+     cat('Pooled covariance: \n')
+     print(cov.pooled)
+     T.sq = t(diff) %%% solve((1/n1+1/n2)*cov.pooled) %%% diff
+     F = TF.convert.twosample(T.sq = T.sq, n1=n1, n2=n2, p=p)
+     p_value = pf(F$F, F$df1, F$df2, lower.tail = FALSE)
+
+     T.sq.alpha = TF.convert.twosample(F=qf(1-alpha,p,n1+n2-p-1),n1=n1,n2=n2,p=p, T_to_F = FALSE)
+     factor = sqrt(1/n1+1/n2)* sqrt(diag(cov.pooled))
+     simontaneous.distance = sqrt(T.sq.alpha$T.sq) * factor
+     simontaneous.CI = cbind(mean_diff - simontaneous.distance, mean_diff + simontaneous.distance)
+     colnames(simontaneous.CI) = c('low', 'high')
+     rownames(simontaneous.CI) = c(1:p)
+     t = qt(1-alpha/2/p, df = n1+n2-2)
+     bonferroni.distance = sqrt(t) * factor
+     bonferroni.CI = cbind(mean_diff - bonferroni.distance, mean_diff + bonferroni.distance)
+     colnames(bonferroni.CI) = c('low', 'high')
+     rownames(bonferroni.CI) = c(1:p)
+     cat('\n-----\nT-square Value: ', T.sq, '\nP-value: ', p_value,'\n-----\n', 'Confident Intervals at ', 100-alpha*100, '% c
onfidnet level:\n',sep = '')
+     linear_combination = solve(cov.pooled) %%% diff
+   }else{
+     cat("Method: Two Population T-square Test with Unequal Variance (Unpooled)\n-----\n")
+     cov.unpooled = 1/n1 * cov1 + 1/n2 *cov2
+     cat('Unpooled covariance: \n')
+     print(cov.unpooled)
+     T.sq = t(diff) %%% solve(cov.unpooled) %%% diff
+     p_value = pchisq(T.sq, p, lower.tail = FALSE)
+
+     chi.sq.alpha = qchisq(1-alpha,p)
+     factor = sqrt(diag(cov.unpooled))
+     simontaneous.distance = sqrt(chi.sq.alpha) * factor
+     simontaneous.CI = cbind(mean_diff - simontaneous.distance, mean_diff + simontaneous.distance)
+     colnames(simontaneous.CI) = c('low', 'high')
+     rownames(simontaneous.CI) = c(1:p)
```

```

+
+   z = qnorm(1-alpha/2/p)
+   bonferroni.distance = sqrt(z) * factor
+   bonferroni.CI = cbind(mean_diff - bonferroni.distance, mean_diff + bonferroni.distance)
+   colnames(bonferroni.CI) = c('low', 'high')
+   rownames(bonferroni.CI) = c(1:p)
+   cat('\n-----\nT-square Value: ', T.sq, '\nP-value: ', p_value, '\n-----\n', 'Confident Intervals at ', 100-alpha*100, '% c
onfident level:\n', sep = '')
+   linear_combination = solve(cov.unpooled) %%% diff
+ }
+ cat('Simontaneous CI: \n')
+ print(simontaneous.CI)
+ cat('\nBonferroni CI: \n')
+ print(bonferroni.CI)
+ cat('\n-----\n')
+ cat('The most critical linear combination to reject H0 is: \n')
+ print(linear_combination)
+ cat('\n-----\n')
+ }

```

2.

```

> TF.convert.twosample = function(T.sq = 0, F=0, n1, n2, p, T_to_F = TRUE){
+   if(T_to_F){
+     F = (n1+n2-p-1)/(n1+n2-2)/p * T.sq
+     df1 = p
+     df2 = n1+n2-p-1
+     returnValue(data.frame(F,df1,df2))
+   }else{
+     T.sq = F*(n1+n2-2)*p/(n1+n2-p-1)
+     df1 = p
+     df2 = n1+n2-2
+     returnValue(data.frame(T.sq,df1,df2))
+   }
+ }

```

3.

```

> MANOVA = function(X,trt){
+   n = as.integer(table(trt))
+   N = length(trt)
+   cat('\n-----\nMANOVA Procedure\n-----\n\nGrand Mean:\n')
+   xbar = apply(X, 2, mean)
+   print(xbar)
+   cat('\nTreatment Means: \n')
+   trtmean = NULL
+   residual = NULL
+   for(i in c(1:dim(X)[2])){
+     k = 1
+     trtmean_i = NULL
+     trtmean_i_long = NULL
+     for(j in c(1:length(n))){
+       mean_ij = mean(X[k:(k+n[j]-1),i])
+       trtmean_i = cbind(trtmean_i, mean_ij)
+       trtmean_i_long = cbind(trtmean_i_long, t(rep(mean_ij,n[j])))
+       k = k+n[j]
+     }
+     trtmean = rbind(trtmean,trtmean_i)
+     residual_i = X[,i] - trtmean_i_long
+     residual = rbind(residual, residual_i)
+   }
+   groups = paste('group', c(1:length(n)))
+   vars = paste('x', c(1:dim(X)[2]), sep = '')
+   colnames(trtmean) = groups
+   rownames(trtmean) = vars
+   print(trtmean)
+
+   cat('\nTreatment Effects: \n')

```

```

+ trteffect = trtmean - xbar
+ print(trteffect)
+ cat('\nResiduals:\n')
+ print(residual)
+ SStrt = 0
+ for(i in c(1:length(n))){
+   SStrt = SStrt + n[i] * trteffect[,i] %% t(trteffect[,i])
+ }
+ SSerror = 0
+ for(i in c(1:dim(residual)[2])){
+   temp = residual[,i]
+   SSerror = SSerror + temp %% t(temp)
+ }
+ SStotal = SStrt + SSerror
+ cat('\n-----\nThe MANOVA Table Elements: \n-----\nSSrt (df = ',length(n)-1, ')\n', sep = '')
+ print(SSrt)
+ cat('\nSSerror (df = ',N-length(n), ')\n', sep = '')
+ print(SSerror)
+ cat('\nSStotal (df = ',N-1, ')\n', sep = '')
+ print(SStotal)
+ wilk = det(SSerror) / det(SStotal)
+ ratio = (1-sqrt(wilk))/sqrt(wilk)
+ correction = -(12-1-(2+3)/2)*log(wilk)
+ cat('\n-----\nwilk\'s Test: \n-----\nlamba = ', wilk, '\ntratio = ', ratio,
+   '\n\tp-value: ',1-pf(ratio, 2*(length(n)-1),2*(N-length(n)-1)),
+   '\nBartlett\'s correction: ', correction, '\n\tp-value: ', 1-pchisq(correction, dim(x)[2]*(length(n)-1)),'\n',sep = '')
+ }

```