

[TYPE THE COMPANY NAME]

STAT 445 Assignment 6

Kun Yang 301178299

Problem 9.18 Do only the maximum likelihood portion of part (d).

```
> (CORR<- matrix(c(1,0.4919,0.2636,0.4653,-0.2277,0.0652,  
+ 0.4919,1,0.3127,0.3506,-0.1917,0.2045,  
+ 0.2635,0.3127,1,0.4108, 0.0647,0.2493,  
+ 0.4653,0.3506,0.4108,1,-0.2249,0.2293,  
+ -0.2277,-0.1917,0.0647,-0.2249,1,-0.2144,  
+ 0.0652,0.2045,0.2493,0.2293,-0.2144,1),ncol=6,byrow=TRUE))
```

```
      [,1] [,2] [,3] [,4] [,5] [,6]  
[1,] 1.0000 0.4919 0.2636 0.4653 -0.2277 0.0652  
[2,] 0.4919 1.0000 0.3127 0.3506 -0.1917 0.2045  
[3,] 0.2635 0.3127 1.0000 0.4108 0.0647 0.2493  
[4,] 0.4653 0.3506 0.4108 1.0000 -0.2249 0.2293  
[5,] -0.2277 -0.1917 0.0647 -0.2249 1.0000 -0.2144  
[6,] 0.0652 0.2045 0.2493 0.2293 -0.2144 1.0000
```

```
> (eig=eigen(CORR))
```

\$values

```
[1] 2.3549250 1.0718567 0.9842486 0.6643858 0.5003897 0.4241942
```

\$vectors

```
      [,1] [,2] [,3] [,4] [,5] [,6]  
[1,] -0.4753543 0.02188910 0.47987482 0.04564932 0.3578888 -0.6425051  
[2,] -0.4719328 -0.01924243 0.20906455 0.70296351 -0.1771358 0.4556533  
[3,] -0.3931540 -0.56061775 -0.26440849 -0.17551515 -0.5973982 -0.2713733  
[4,] -0.4963538 -0.07723024 0.03226116 -0.60426514 0.3238962 0.5260914  
[5,] 0.2563177 -0.80502150 0.01294351 0.21817123 0.4823355 0.0768220  
[6,] -0.2910014 0.17559671 -0.80925398 0.24539324 0.3822272 -0.1524794
```

###analysis with 3 factors

```
> (FA=factanal(covmat=CORR,factors=3) )
```

Call:

```
factanal(factors = 3, covmat = CORR)
```

Uniquenesses:

```
[1] 0.005 0.666 0.005 0.607 0.683 0.720
```

Loadings:

```
      Factor1 Factor2 Factor3  
[1,] 0.994  
[2,] 0.466 0.211 0.268  
[3,] 0.201 0.976  
[4,] 0.429 0.316 0.330  
[5,] -0.208 0.134 -0.505  
[6,]      0.227 0.478
```

```
      Factor1 Factor2 Factor3  
SS loadings      1.474 1.170 0.670  
Proportion Var   0.246 0.195 0.112  
Cumulative Var   0.246 0.441 0.552
```

The degrees of freedom for the model is 0 and the fit was 0.0016

Handwritten red notes:
A checkmark is drawn next to the 'Call:' line.
A red box is drawn around the 'Uniquenesses:' line.
A red circle is drawn around the 'Uniquenesses:' line, with the text '19/20' written inside it.

```

> str(FA)
List of 11
 $ converged   : logi TRUE
 $ loadings    : loadings [1:6, 1:3] 0.994 0.466 0.201 0.429 -0.208 ...
 .. attr(*, "dimnames")=List of 2
 .. ..$ : NULL
 .. ..$ : chr [1:3] "Factor1" "Factor2" "Factor3"
 $ uniquenesses: num [1:6] 0.005 0.666 0.005 0.607 0.683 ...
 $ correlation : num [1:6, 1:6] 1 0.492 0.264 0.465 -0.228 ...
 $ criteria     : Named num [1:3] 0.00157 18 18
 .. attr(*, "names")= chr [1:3] "objective" "counts.function" "counts.gradient"
 $ factors      : num 3
 $ dof          : num 0
 $ method       : chr "mle"
 $ rotmat       : num [1:3, 1:3] 0.7539 -0.6544 0.0577 0.6533 0.7561 ...
 $ n.obs        : logi NA
 $ call         : language factanal(factors = 3, covmat = CORR)
 - attr(*, "class")= chr "factanal"

```

> ###communalities

```
> L=FA$loadings
```

```
> (hi2=(L%*%t(L)))
```

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]
[1,]	0.9950001	0.4918611	0.26350000	0.4653067	-0.22769436	0.0652224
[2,]	0.4918611	0.3336753	0.31273101	0.3551980	-0.20419524	0.1870385
[3,]	0.2635000	0.3127310	0.99500005	0.4107643	0.06469249	0.2493014
[4,]	0.4653067	0.3551980	0.41076428	0.3929169	-0.21361990	0.2394891
[5,]	-0.2276944	-0.2041952	0.06469249	-0.2136199	0.31692365	-0.2160465
[6,]	0.0652224	0.1870385	0.24930139	0.2394891	-0.21604647	0.2804297

> ###specific variances

```
> (psi=diag(FA$uniquenesses))
```

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]
[1,]	0.005	0.0000000	0.000	0.0000000	0.0000000	0.0000000
[2,]	0.000	0.6663258	0.000	0.0000000	0.0000000	0.0000000
[3,]	0.000	0.0000000	0.005	0.0000000	0.0000000	0.0000000
[4,]	0.000	0.0000000	0.000	0.6070835	0.0000000	0.0000000
[5,]	0.000	0.0000000	0.000	0.0000000	0.683076	0.0000000
[6,]	0.000	0.0000000	0.000	0.0000000	0.0000000	0.7195693

> ###residual matrix and round it to 4 decimal places

```
> RezMat=CORR-(hi2+psi)
```

```
> round(RezMat,4)
```

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]
[1,]	0	0.0000	1e-04	0.0000	0.0000	0.0000
[2,]	0	0.0000	0e+00	-0.0046	0.0125	0.0175
[3,]	0	0.0000	0e+00	0.0000	0.0000	0.0000
[4,]	0	-0.0046	0e+00	0.0000	-0.0113	-0.0102
[5,]	0	0.0125	0e+00	-0.0113	0.0000	0.0016

Agg-18

[6,] 0 0.0175 0e+00 -0.0102 0.0016 0.0000 ✓

Diagonal elements of the residual matrix are 0 which means that (hi2+psi) is close to the correlation matrix. So $m = 3$ factor solution represents the observed correlations quite well.

Factor 1 puts significant and maximum weight on bluegill(x1). Black crappie, smallmouth bass, Largemouth bass, Walleye, Northern pike have significantly smaller weights than the weights on the bluegill group. The first factor emphasize on bluegill.

description of factor 1 is grouping those loadings with largest positive weights (not just one weight)

Factor 2 puts significant and maximum weight on smallmouth bass (x3). Bluegill, Black crappie, Largemouth bass, Walleye, Northern pike have significantly smaller weights than the weights on the smallmouth bass group. So the second factor emphasize on smallmouth bass.

grouping!

Factor 3 is negatively driven by walleye. But it is not very significant.

yes but you have to force the group with largest positive weights.

These three factors collectively accounted for 55.2% of total variation. It represents quite small proportion of variation. But if we conduct the analysis by $m=2$, the cumulative variation will be even lower and the second factor is hard to interpret meaningfully. Therefore, I choose $m=3$.

You could also look at the p-value.

Analysis with 2 factors as following:

```
> (FA=factanal(covmat=CORR,factors=2) )
```

Call:

```
factanal(factors = 2, covmat = CORR)
```

Uniquenesses:

```
[1] 0.453 0.600 0.005 0.581 0.829 0.914
```

Loadings:

	Factor1	Factor2
[1,]	0.279	0.685
[2,]	0.326	0.542
[3,]	0.997	
[4,]	0.423	0.491
[5,]		-0.410
[6,]	0.253	0.148

hard to interpret.

	Factor1	Factor2
SS loadings	1.424	1.195
Proportion Var	0.237	0.199
Cumulative Var	0.237	0.437

The degrees of freedom for the model is 4 and the fit was 0.0935

Problem 9.19

Part (a) Obtain only the maximum likelihood estimates for the factor analysis model with $m=2$ & $m=3$.

```
> Data.matrix<-read.csv("Salespeopledata.csv")
```

```
> (CORR=cor(Data.matrix))
```

	SG	SP	NA.	CT	MR	AR	MT
SG	1.0000000	0.9260758	0.8840023	0.5720363	0.7080738	0.6744073	0.9273116
SP	0.9260758	1.0000000	0.8425232	0.5415080	0.7459097	0.4653880	0.9442960
NA.	0.8840023	0.8425232	1.0000000	0.7003630	0.6374712	0.6410886	0.8525682
CT	0.5720363	0.5415080	0.7003630	1.0000000	0.5907360	0.1469074	0.4126395
MR	0.7080738	0.7459097	0.6374712	0.5907360	1.0000000	0.3859502	0.5745533
AR	0.6744073	0.4653880	0.6410886	0.1469074	0.3859502	1.0000000	0.5663721
MT	0.9273116	0.9442960	0.8525682	0.4126395	0.5745533	0.5663721	1.0000000

```
> (eig=eigen(CORR))
```

\$values

```
[1] 5.03459779 0.93351614 0.49791975 0.42124549 0.08104043 0.02034063 0.01133977
```

\$vectors

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]
[1,]	-0.4336719	0.111754422	0.075488541	-0.04237344	0.6324942624	0.3365963	0.52782527
[2,]	-0.4202136	-0.029287495	0.442478953	0.01075255	-0.0001182093	-0.7853424	0.09948330
[3,]	-0.4210510	-0.009201975	-0.204189315	-0.32492838	-0.7010262539	0.1568114	0.39916419
[4,]	-0.2942863	-0.668415809	-0.451492333	-0.30271208	0.2610080204	-0.1141710	-0.29995962
[5,]	-0.3490920	-0.294944379	-0.005921773	0.84660356	-0.1742634819	0.1969091	-0.07231139
[6,]	-0.2891669	0.642377957	-0.603779622	0.15367411	0.0869586057	-0.2362610	-0.22844351
[7,]	-0.4074041	0.200367651	0.434039576	-0.24601320	-0.0495826418	0.3711105	-0.63622351

###analysis with 3 factors (m=3)

```
> (FA=factanal(covmat=CORR,factors=3) )
```

Call:

```
factanal(factors = 3, covmat = CORR)
```

Uniquenesses:

	SG	SP	NA.	CT	MR	AR	MT
	0.039	0.034	0.088	0.005	0.447	0.005	0.038

Loadings:

	Factor1	Factor2	Factor3
SG	0.793	0.374	0.438
SP	0.911	0.317	0.185
NA.	0.651	0.544	0.438
CT	0.255	0.964	
MR	0.542	0.465	0.207
AR	0.299		0.950
MT	0.917	0.180	0.298

	Factor1	Factor2	Factor3
SS loadings	3.175	1.718	1.453
Proportion Var	0.454	0.245	0.208
Cumulative Var	0.454	0.699	0.906

The degrees of freedom for the model is 3 and the fit was 1.4186

```

> str(FA)
List of 11
 $ converged   : logi TRUE
 $ loadings    : loadings [1:7, 1:3] 0.793 0.911 0.651 0.255 0.542 ...
  .. attr(*, "dimnames")=List of 2
  .. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
  .. ..$ : chr [1:3] "Factor1" "Factor2" "Factor3"
 $ uniquenesses: Named num [1:7] 0.0386 0.0345 0.0881 0.005 0.4466 ...
  .. attr(*, "names")= chr [1:7] "SG" "SP" "NA." "CT" ...
 $ correlation : num [1:7, 1:7] 1 0.926 0.884 0.572 0.708 ...
  .. attr(*, "dimnames")=List of 2
  .. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
  .. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
 $ criteria    : Named num [1:3] 1.42 27 27
  .. attr(*, "names")= chr [1:3] "objective" "counts.function" "counts.gradient"
 $ factors     : num 3
 $ dof         : num 3
 $ method      : chr "mle"
 $ rotmat      : num [1:3, 1:3] 0.612 0.714 0.341 -0.6 0.699 ...
 $ n.obs       : logi NA
 $ call        : language factanal(factors = 3, covmat = CORR)
 - attr(*, "class")= chr "factanal"

```

###communalities

```

> L=FA$loadings
> (hi2=(L%*%t(L)))

```

	SG	SP	NA.	CT	MR	AR	MT
SG	0.9614284	0.9228135	0.9120898	0.5714373	0.6949357	0.6738835	0.9255320
SP	0.9228135	0.9655192	0.8471023	0.5417813	0.6799465	0.4654561	0.9481930
NA.	0.9120898	0.8471023	0.9118756	0.6991264	0.6969691	0.6402871	0.8255826
CT	0.5714373	0.5417813	0.6991264	0.9950434	0.5910466	0.1469508	0.4130097
MR	0.6949357	0.6799465	0.6969691	0.5910466	0.5533820	0.3841948	0.6425648
AR	0.6738835	0.4654561	0.6402871	0.1469508	0.3841948	0.9950317	0.5669006
MT	0.9255320	0.9481930	0.8255826	0.4130097	0.6425648	0.5669006	0.9624901

###specific variances

```

> (psi=diag(FA$uniquenesses))

```

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]
[1,]	0.03857165	0.00000000	0.00000000	0.000	0.00000000	0.000	0.00000000
[2,]	0.00000000	0.03448071	0.00000000	0.000	0.00000000	0.000	0.00000000
[3,]	0.00000000	0.00000000	0.08812176	0.000	0.00000000	0.000	0.00000000
[4,]	0.00000000	0.00000000	0.00000000	0.005	0.00000000	0.000	0.00000000
[5,]	0.00000000	0.00000000	0.00000000	0.000	0.4466205	0.000	0.00000000
[6,]	0.00000000	0.00000000	0.00000000	0.000	0.00000000	0.005	0.00000000
[7,]	0.00000000	0.00000000	0.00000000	0.000	0.00000000	0.000	0.0375098

###residual matrix and round it to 4 decimal places

```
> RezMat=CORR-(hi2+psi)
```

```
> round(RezMat,4)
```

	SG	SP	NA.	CT	MR	AR	MT
SG	<u>0.0000</u>	0.0033	-0.0281	0.0006	0.0131	0.0005	0.0018
SP	0.0033	<u>0.0000</u>	-0.0046	-0.0003	0.0660	-0.0001	-0.0039
NA.	-0.0281	-0.0046	<u>0.0000</u>	0.0012	-0.0595	0.0008	0.0270
CT	0.0006	-0.0003	0.0012	<u>0.0000</u>	-0.0003	0.0000	-0.0004
MR	0.0131	0.0660	-0.0595	-0.0003	<u>0.0000</u>	0.0018	-0.0680
AR	0.0005	-0.0001	0.0008	0.0000	0.0018	<u>0.0000</u>	-0.0005
MT	0.0018	-0.0039	0.0270	-0.0004	-0.0680	-0.0005	<u>0.0000</u>

Diagonal elements of the residual matrix are 0 which means that (hi2+psi) is close to the correlation matrix. So $m = 3$ factor solution represents the observed correlations quite well.

###analysis with 2 factors

```
> (FA=factanal(covmat=CORR,factors=2) )
```

Call:

```
factanal(factors = 2, covmat = CORR)
```

Uniquenesses:

	SG	SP	NA.	CT	MR	AR	MT
	0.069	0.070	0.123	0.005	0.474	0.614	0.029

Loadings:

	Factor1	Factor2
SG	0.852	0.452
SP	0.868	0.419
NA.	0.717	0.602
CT	0.148	0.987
MR	0.501	0.525
AR	0.619	
MT	0.946	0.277

	Factor1	Factor2
SS loadings	3.545	2.071
Proportion Var	0.506	0.296
Cumulative Var	0.506	0.802

The degrees of freedom for the model is 8 and the fit was 2.6337

```
> str(FA)
```

List of 11

```
$ converged : logi TRUE
$ loadings : loadings [1:7, 1:2] 0.852 0.868 0.717 0.148 0.501 ...
..- attr(*, "dimnames")=List of 2
.. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
.. ..$ : chr [1:2] "Factor1" "Factor2"
$ uniquenesses: Named num [1:7] 0.0692 0.0704 0.1233 0.005 0.4736 ...
..- attr(*, "names")= chr [1:7] "SG" "SP" "NA." "CT" ...
$ correlation : num [1:7, 1:7] 1 0.926 0.884 0.572 0.708 ...
..- attr(*, "dimnames")=List of 2
.. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
```

Why do you
print this?
and why
many times


```

.. ..$ : chr [1:7] "SG" "SP" "NA." "CT" ...
$ criteria : Named num [1:3] 2.63 33 33
..- attr(*, "names")= chr [1:3] "objective" "counts.function" "counts.gradient"
$ factors : num 2
$ dof : num 8
$ method : chr "mle"
$ rotmat : num [1:2, 1:2] 0.95 -0.312 0.312 0.95
$ n.obs : logi NA
$ call : language factanal(factors = 2, covmat = CORR)
- attr(*, "class")= chr "factanal"

```

###communalities

```

> L=FA$loadings
> (hi2=(L%*%t(L)))

```

	SG	SP	NA.	CT	MR	AR	MT
SG	0.9308083	0.9295188	0.8834712	0.5720627	0.6642317	0.5543372	0.9311883
SP	0.9295188	0.9296182	0.8749729	0.5413952	0.6547850	0.5624008	0.9373111
NA.	0.8834712	0.8749729	0.8766896	0.6996404	0.6751663	0.4798309	0.8449575
CT	0.5720627	0.5413952	0.6996404	0.9950121	0.5918666	0.1504777	0.4126431
MR	0.6642317	0.6547850	0.6751663	0.5918666	0.5264121	0.3412919	0.6189370
AR	0.5543372	0.5624008	0.4798309	0.1504777	0.3412919	0.3863622	0.6017601
MT	0.9311883	0.9373111	0.8449575	0.4126431	0.6189370	0.6017601	0.9711829

###specific variances

```

> (psi=diag(FA$uniquenesses))
      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]      [,7]
[1,] 0.0691916 0.0000000 0.0000000 0.000 0.0000000 0.0000000 0.0000000
[2,] 0.0000000 0.07038038 0.0000000 0.000 0.0000000 0.0000000 0.0000000
[3,] 0.0000000 0.0000000 0.1233088 0.000 0.0000000 0.0000000 0.0000000
[4,] 0.0000000 0.0000000 0.0000000 0.005 0.0000000 0.0000000 0.0000000
[5,] 0.0000000 0.0000000 0.0000000 0.000 0.4735849 0.0000000 0.0000000
[6,] 0.0000000 0.0000000 0.0000000 0.000 0.0000000 0.6136386 0.0000000
[7,] 0.0000000 0.0000000 0.0000000 0.000 0.0000000 0.0000000 0.02881701

```

###residual matrix and round it to 4 decimal places

```

> RezMat=CORR-(hi2+psi)
> round(RezMat,4)
      SG      SP      NA.      CT      MR      AR      MT
SG  0.0000 -0.0034  0.0005  0.0000  0.0438  0.1201 -0.0039
SP -0.0034  0.0000 -0.0324  0.0001  0.0911 -0.0970  0.0070
NA. 0.0005 -0.0324  0.0000  0.0007 -0.0377  0.1613  0.0076
CT  0.0000  0.0001  0.0007  0.0000 -0.0011 -0.0036  0.0000
MR  0.0438  0.0911 -0.0377 -0.0011  0.0000  0.0447 -0.0444
AR  0.1201 -0.0970  0.1613 -0.0036  0.0447  0.0000 -0.0354
MT -0.0039  0.0070  0.0076  0.0000 -0.0444 -0.0354  0.0000

```

Diagonal elements of the residual matrix are 0 which means that (hi2+psi) is close to the correlation matrix, so m = 2 factor solution represents the observed correlations quite well.

Part (b) Interpret these estimates where possible.

For analysis with 3 factors (m=3)

Factor 1 puts maximum weight on mathematics(x7) test and it places large weights on sales growth (x1) and profitability of sales (x2) and new account sales(x3). Thus, the first factor can be described as the salesperson's performance (sales growth, profitability of sales, new account sales) with one's mathematical ability.

Factor 2 puts significant and maximum weight on creativity test. Thus, the second factor can be described as creative ability.

Factor 3 puts significant and maximum weight on abstract reasoning test. Thus, the third factor can be described as abstract reasoning ability

These three factors collectively accounted for 90.6% of total variation.

For analysis with 2 factors (m=2)

Factor 1 puts maximum weight on mathematics(x7) test and it places large weights on sales growth (x1) and profitability of sales (x2) and new account sales(x3). Thus, the first factor can be described as the salesperson's performance (sales growth, profitability of sales, new account sales) with one's mathematical ability.

Factor 2 puts significant and maximum weight on creativity test. Thus, the second factor can be described as creative ability.

These three factors collectively accounted for 80.2% of total variation.

Part (d) As stated. I.e., perform the specified tests and, in light of the results of these tests and the interpretations in Part (b), draw an appropriate conclusion on the best value for m.

$H_0: \Sigma = L'L + D_\psi$, with $m = 3$, at level $\alpha = 0.01$

For analysis with 3 factors (m=3)

```
> (A<-det(hi2+psi))  $\Rightarrow L'L + \psi$   
[1] 7.727395e-05  
> (B<-det(CORR))  
[1] 1.842687e-05 =  $S_n$   
> (C<-50-1-(2*7+4*3+5)/6)  
[1] 43.83333  
> (Test<-C*log(A/B))  
[1] 62.83714
```

$$\left(50 - 1 - \frac{2*7+4*3+5}{6}\right) \ln\left(\frac{|L'L+D_\psi|}{|S_n|}\right)$$

$$v = \frac{1}{2}[(7-3)^2 - 7 - 3] = 3$$

$$\chi^2 = 11.34 \quad \text{with} \quad df=3 \quad \alpha=0.01$$

So we reject H_0 for $m=3$.

✓ you could use p-values
returned from
factanal f.m.

For analysis with 2 factors (m=2)

$H_0: \Sigma = L'L + D_\psi$, with $m = 2$, at level $\alpha = 0.01$

> (A<-det(hi2+psi)) $\approx L'L + \psi$

[1] 0.0002572145

> (B<-det(CORR)) $= S_n$

[1] 1.842687e-05

> (C<-50-1-(2*7+4*2+5)/6) -

[1] 44.5

> (Test<-C*log(A/B))

[1] 117.3065

$$\left(50 - 1 - \frac{2 \cdot 7 + 4 \cdot 2 + 5}{6}\right) \ln\left(\frac{|L'L + D_\psi|}{|S_n|}\right)$$

$$v = \frac{1}{2}[(7-2)^2 - 7 - 2] = 8$$

$$\chi^2 = 20.09 \quad \text{with} \quad \text{df} = 8 \quad \alpha = 0.01$$

So we reject H_0 for $m=2$.

Therefore, we can say neither of the $m = 2$, $m = 3$ models can fit the χ^2 criterion.

But p-value are closer to α when $m=3$. And in light of part (b), I prefer $m=3$, because all of the 3 factors can be interpreted meaningfully. When $m=3$, diagonal elements of the residual matrix are 0. These three factors collectively accounted for 90.6% of total variation.

Hello Biljana,

I have a question,
Do we need to include str(LFA)
in the factor analysis every time?
How important it is?

Thank you for your answer !!!
Kun Yung.