# HUDM6122 Homework\_05

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# 0.1 Github Address

All my latest homework can be found on Github:  $https://github.com/cgpan/hudm6122\_homeworks$ . Thanks for checking if interested.

# 0.2 Ex 5.1

Show how the result rises from the assumptions of uncorrelated factors, independence of the specific variates, and independence of common factors and specific variances. What form does take if the factors are allowed to be correlated?

#### MY SOLUTION:

Based on the assumption of Exploratory Factor Analysis (EFA), a set of observed variables  $\mathbf{x}$  assumed to be linked to a set of latent variables  $\mathbf{f}$ . Therefore, we can have a regression model in matrix form

$$x = \Lambda f + u$$

, where  $\Lambda$  is a  $q \times k$  matrix of factor loadings (a.k.a., the coefficients of the regression model), and the u is the vector of unexplained error of each observed variables.

Let's take the variance of the formula above

$$V(\boldsymbol{x}) = V(\boldsymbol{\Lambda}\boldsymbol{f} + \boldsymbol{u})$$

. Based on the operation rule of variance, like

$$V(a+b) = V(a) + V(b) + 2Cov(ab)$$

, we combined the two formulas above, then

$$V(\boldsymbol{x}) = V(\boldsymbol{\Lambda}\boldsymbol{f} + \boldsymbol{u}) = V(\boldsymbol{\Lambda}\boldsymbol{f}) + V(\boldsymbol{u}) + 2Cov(\boldsymbol{\Lambda}\boldsymbol{f}\boldsymbol{u})$$

. Since the we assumed that the error terms are uncorrelated with the factors, therefore the  $Cov(\mathbf{\Lambda} \mathbf{f} \mathbf{u}) = 0$ . Then, we can continue to drive the variance formula as

$$V(\boldsymbol{x}) = V(\boldsymbol{\Lambda}\boldsymbol{f}) + V(\boldsymbol{u}) = \boldsymbol{\Lambda}V(\boldsymbol{f})\boldsymbol{\Lambda}^T + \Psi$$

. In addition, we assumed that the factors are uncorrelated with each other. The V(f) is actually an identity matrix. Therefore, the formula can be written as

$$V(\boldsymbol{x}) = \boldsymbol{\Lambda} V(\boldsymbol{f}) \boldsymbol{\Lambda}^T + \boldsymbol{\Psi} = \boldsymbol{\Lambda} \boldsymbol{\Lambda}^T + \boldsymbol{\Psi}$$

. Finally, the formula can be written as

$$\Sigma = \Lambda \Lambda^T + \Psi$$

.

If we allow the factors to be correlated with each other, then the V(f) is not an identity matrix. Let's use the greek letter  $\Phi$  to represent the variance matrix of loadings f. Thus, the formula should be

$$\mathbf{\Sigma} = \mathbf{\Lambda} \mathbf{\Phi} \mathbf{\Lambda}^T + \mathbf{\Psi}$$

.

### 0.3 Ex 5.2

Show that the communalities in a factor analysis model are unaffected by the transformation . . .

#### MY SOLUTION:

This question mentioned that we need to use the transformed factor loadings  $\Lambda^* = \Lambda M$ . Let's assume that M is an  $k \times k$  orthogonal matrix. We can re-write the basic regression equation linking the observed and the factors as:

$$\boldsymbol{x} = (\boldsymbol{\Lambda} \boldsymbol{M})(\boldsymbol{M}^T \boldsymbol{f}) + \boldsymbol{u}$$

.

Using the rule of variance, we can have

$$\boldsymbol{\Sigma} = (\boldsymbol{\Lambda}\boldsymbol{M})(\boldsymbol{\Lambda}\boldsymbol{M})^T + \boldsymbol{\Psi}$$

. Since the M is a orthogonal matrix and  $MM^T = I$ . Therefore, the variance equation can be written as

$$\mathbf{\Sigma} = \mathbf{\Lambda} \mathbf{\Lambda}^T + \Psi$$

. That is, the transformed factor loadings  $\mathbf{\Lambda}^* = \mathbf{\Lambda} \mathbf{M}$  will not influence the communalities (i.e.,  $\mathbf{\Lambda} \mathbf{\Lambda}^T$ ) in the a factor analysis model.

# 0.4 Ex 5.3

Give a formula for the proportion of variance explained by the jth factor estimated by the principal factor approach.

### MY SOLUTION:

The proportion of variance explained by the jth factor represents the proportion of the total variance in the observed variables that is accounted for by that factor alone. Therefore, the formula could be

$$Proportion_j = \frac{\sum_{i=1}^{q} \lambda_{ij}^2}{\mathbf{\Lambda} \mathbf{\Lambda}^T}$$

.

#### 0.5 Ex 5.4

Apply the factor analysis model separately to the life expectancies of men and women and compare the results.

#### MY SOLUTION:

The textbook does not provide the original dataset. Based on the code in the MVA, I create the dataset via a separated r file named "HW05 Test". This file created the life.rdata and life.csv dataset in the same file folder.

```
> load("life.rdata")
> head(life)
          m0 m25 m50 m75 w0 w25 w50 w75
Algeria
          63 51 30 13 67 54 34
Cameroon
          34 29 13
                       5 38
                             32 17
                                       6
Madagascar 38 30
                  17
                       7 38
                             34
                                 20
                                       7
Mauritius 59 42 20
                       6 64
                             46 25
                                      8
Reunion
          56 38 18
                       7 62
                             46 25 10
Seychelles 62 44 24
                       7 69 50 28 14
> # subset the male and female dataset
> life_male <- life[,1:4]</pre>
> life_female <- life[,5:8]</pre>
> # test the number of factors needed for the male and female dataset separately
> sapply(1, function(f)
   factanal(life_male, factors=f, method="mle")$PVAL)
   objective
0.0007284301
> sapply(1, function(f)
   factanal(life_female, factors=f, method="mle")$PVAL)
   objective
4.738464e-12
```

When test the number of the factors from 1 to larger number, there is always a warning that N factors are too many for N variables. More details can be found on Page 143 of the textbook or here https://stats.stackexchange.com/questions/593452/efa-n-factors-are-too-many-for-n-variables

The results suggest that an one-factor solution might be adequate to account for the observed covariances in the data.

Next, I run the one-factor solution for both male and female datasets.

```
> factanal(life_male, factors = 1, method="mle")
Call:
factanal(x = life_male, factors = 1, method = "mle")
Uniquenesses:
       m25
  mΟ
             m50 m75
0.594 0.552 0.005 0.434
Loadings:
   Factor1
m0 0.638
m25 0.669
m50 0.998
m75 0.752
               Factor1
                 2.415
SS loadings
Proportion Var
                 0.604
Test of the hypothesis that 1 factor is sufficient.
```

```
The chi square statistic is 14.45 on 2 degrees of freedom.
The p-value is 0.000728
> factanal(life_female, factors = 1, method="mle")
factanal(x = life_female, factors = 1, method = "mle")
Uniquenesses:
       w25
             w50
                    w75
  wO
0.220 0.005 0.115 0.526
Loadings:
   Factor1
w0 0.883
w25 0.998
w50 0.941
w75 0.689
               Factor1
                 3.134
SS loadings
                 0.784
Proportion Var
Test of the hypothesis that 1 factor is sufficient.
The chi square statistic is 52.15 on 2 degrees of freedom.
The p-value is 4.74e-12
```

#### 0.6 Ex 5.6

The matrix below shows the correlations between ratings on nine statements about pain made by 123 people suffering from extreme pain. Each statement was scored on a scale from 1 to 6, ranging from agreement to disagreement. The nine pain statements were as follows:

#### MY SOLUTION:

First, to change the lower triangular matrix into the complete correlation matrix.

The correlation matrix looks good. Next, I run the PCA first.

```
> # run the PCA first
> # use prcomp to calculate the principal components
> pca <- prcomp(corr_symmetric, scale. = FALSE)</pre>
> # get the PCA results
> summary(pca)
Importance of components:
                          PC1
                                 PC2
                                         PC3
                                                 PC4
                                                        PC5
                                                                PC6
                                                                         PC7
Standard deviation
                       1.1594 0.5014 0.31822 0.18698 0.1734 0.15725 0.09446
Proportion of Variance 0.7466 0.1396 0.05624 0.01942 0.0167 0.01373 0.00496
Cumulative Proportion 0.7466 0.8863 0.94250 0.96192 0.9786 0.99235 0.99730
                           PC8
                                     PC9
Standard deviation
                       0.06968 3.631e-17
Proportion of Variance 0.00270 0.000e+00
Cumulative Proportion 1.00000 1.000e+00
> # draw the scree plot
> plot(pca, type = "1",
+ main = "Scree Plot")
```

# **Scree Plot**

