Factor Analysis

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STAT 4690-Applied Multivariate Analysis

Latent variable models

- With PCA, we saw how we could reduce the dimension of data using the eigenvectors of the sample covariance matrix.
- Conversely, we could construe PCA has a generative model, where the principal components give rise to the observed data.
- Latent Variable Models formalise this idea:
 - Latent (i.e. unobserved) variables F give rise to observed data Y through a specified model.

Factor Analysis i

- Factor Analysis is a special kind of latent variable model.
- Let Y be a p-dimensional vector with mean μ and covariance matrix Σ .
- Let **F** be a *m*-dimensional *latent* vector.
- The *orthogonal factor model* is given by

$$\mathbf{Y} - \mu = L\mathbf{F} + \mathbf{E},$$

where L is a $p \times m$ matrix of factor loadings, and \mathbf{E} is a p-dimensional vector of errors.

Factor Analysis ii

- F are also called *common factors*; E are also called *specific factors*.
- **Note**: This is essentially a multivariate regression model, but where the covariates are unobserved.

Assumptions i

- The model above is generally not identifiable, since there are too many parameters.
- We therefore need to impose further restrictions:
 - $E(\mathbf{F}) = 0$
 - $Cov(\mathbf{F}) = I$
 - $E(\mathbf{E}) = 0$

$$\mathbf{Cov}(\mathbf{E}) = \Psi = \begin{pmatrix} \psi_1 & 0 & \cdots & 0 \\ 0 & \psi_2 & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & \psi_p \end{pmatrix}$$

$$\mathbf{Cov}(\mathbf{F}, \mathbf{E}) = 0$$

Assumptions ii

- In other words:
- Both common and specific factors have mean zero;
- They are uncorrelated;
- The common factors are mutually uncorrelated and standardised;
- The specific factors each affect only one observed variable.

Structured Covariance i

• As a consequence of these assumptions, we can derive an assumption on the structure of $\Sigma = \text{Cov}(\mathbf{Y})$:

$$\Sigma = E\left((\mathbf{Y} - \mu)(\mathbf{Y} - \mu)^{T}\right)$$

$$= E\left((L\mathbf{F} + \mathbf{E})(L\mathbf{F} + \mathbf{E})^{T}\right)$$

$$= LE(\mathbf{F}\mathbf{F}^{T})L + E(\mathbf{E}\mathbf{F}^{T})L^{T} + LE(\mathbf{F}\mathbf{E}^{T}) + E(\mathbf{E}\mathbf{E}^{T})$$

$$= LIL^{T} + 0L^{T} + L0 + \Psi$$

$$= LL^{T} + \Psi.$$

Structured Covariance ii

Similarly, we can show that

$$Cov(\mathbf{Y}, \mathbf{F}) = L.$$

• If we write ℓ_{ij} for the (i,j)-th element of L, we see that

$$Var(Y_i) = \sum_{k=1}^{m} \ell_{ik}^2 + \psi_i.$$

 Crucially, these assumptions are testable. In other words, we can check whether they are reasonable for our data.

Example i

- Let's look at an example where there is no solution.
- Assume p = 3, m = 1, with

$$\Sigma = \begin{pmatrix} 1 & 0.9 & 0.7 \\ 0.9 & 1 & 0.4 \\ 0.7 & 0.4 & 1 \end{pmatrix}.$$

 From our assumptions on the covariance structure, we derive a system of equations

$$1 = \ell_{11}^2 + \psi_1 \quad 0.9 = \ell_{11}\ell_{21} \quad 0.7 = \ell_{11}\ell_{31}$$
$$1 = \ell_{22}^2 + \psi_2 \quad 0.4 = \ell_{21}\ell_{31}$$
$$1 = \ell_{33}^2 + \psi_3$$

Example ii

• From $0.7 = \ell_{11}\ell_{31}$ and $0.4 = \ell_{21}\ell_{31}$, we get

$$\ell_{21} = \frac{0.4}{0.7} \ell_{11}.$$

• But since $0.9 = \ell_{11}\ell_{21}$, we can conclude that

$$\ell_{11} = \pm 1.255.$$

- However, since the first component Y_1 has unit variance, $\ell_{11} = \operatorname{Corr}(Y_1, F_1)$, and therefore the correlation is out of bounds.
- Similarly, we get

$$\psi_1 = 1 - \ell_{11}^2 = 1 - 1.575 = -0.575.$$

Example iii

• But since ψ_1 is the variance of the first error term, we once again get a non-sensical solution.

Factor Rotation i

- Even with our assumptions above, our model is still not uniquely identified.
- Let T be an $m \times m$ orthogonal matrix. We have

$$\mathbf{Y} - \mu = L\mathbf{F} + \mathbf{E}$$
$$= LTT^{T}\mathbf{F} + \mathbf{E}$$
$$= \tilde{L}\tilde{\mathbf{F}} + \mathbf{E},$$

where $\tilde{L} = LT$ and $\tilde{\mathbf{F}} = T^T\mathbf{F}$.

Factor Rotation ii

Both models lead to the same covariance matrix:

$$\Sigma = LL^T + \Psi = LTT^TL^T + \Psi = \tilde{L}\tilde{L}^T + \Psi.$$

- As we will see, this will turn out to be a blessing in disguise:
 - We will impose a uniqueness condition to get one solution.
 - Then we will rotate our solution using T to improve interpretation.

Estimation-Principal Component Method i

 Recall the spectral decomposition of the covariance matrix:

$$\Sigma = \sum_{i=1}^{p} \lambda_i w_i w_i^T,$$

with $\lambda_1 \geq \cdots \geq \lambda_p$.

• If we let W be the matrix whose i-th column is $\sqrt{\lambda_i}w_i$, we can rewrite the spectral decomposition as

$$\Sigma = WW^T.$$

• In other words, if we let m=p and $\Psi=0$, we see that we recover the orthogonal factor model with L=W.

Estimation-Principal Component Method ii

- Of course, this is not very satisfactory, as the dimension of the common factors is the same as that of the original data.
- Instead, we select m < p using one of the methods we discussed with PCA and we approximate

$$\Sigma \approx \sum_{i=1}^{m} \lambda_i w_i w_i^T.$$

• If we let L be the $p \times m$ matrix whose i-th column is $\sqrt{\lambda_i}w_i$, we can estimate Ψ as follows:

$$\psi_i = \sigma_{ii} - \sum_{j=1}^m \ell_{ij}^2.$$

Estimation-Principal Component Method iii

Algorithm

- 1. Let $\hat{\lambda}_1 \cdots > \hat{\lambda}_p$ and $\hat{w}_1, \ldots, \hat{w}_p$ be the eigenvalues and eigenvectors of the covariance matrix S_n .
- 2. Select m using one of the PCA criteria.
- 3. Estimate \hat{L} with the $p \times m$ matrix whose i-th column is $\sqrt{\hat{\lambda}_i \hat{w}_i}$.
- 4. Estimate $\hat{\Psi}$ with the diagonal elements of $S_n \hat{L}\hat{L}^T$.

Example i

```
library(psych)
dim(bfi)
## [1] 2800 28
names(bfi)
```

Example ii

```
##
    [1] "A1"
                       "A2"
                                    "A3"
                                                  "A4"
    [6] "C1"
                       "C2"
                                    "C3"
                                                  "C4"
##
## [11] "E1"
                       "E2"
                                    "F.3"
                                                  "F.4"
## [16] "N1"
                       "N2"
                                    "N3"
                                                  "N4"
                                    "03"
                                                  "04"
## [21] "01"
                       "02"
## [26] "gender"
                       "education" "age"
```

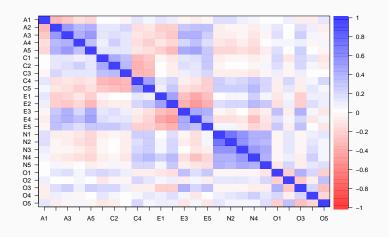
```
library(tidyverse)

data <- bfi %>%
  select(-gender, -education, -age) %>%
  filter(complete.cases(.))
```

Example iii

cor.plot(cor(data))

Example iv



Example v

```
decomp <- prcomp(data)
summary(decomp)$importance[,1:3]</pre>
```

```
## PC1 PC2 PC3
## Standard deviation 3.291635 2.451538 2.030393
## Proportion of Variance 0.215650 0.119620 0.082050
## Cumulative Proportion 0.215650 0.335270 0.417320
```

Example vi

```
cum_prop <- decomp %>%
  summary %>%
  .[["importance"]] %>%
  .["Cumulative Proportion",]

which(cum_prop > 0.8)[1]
```

```
## PC14
## 14
```

Example vii

```
Lhat <- decomp$rotation[,1:14] %*%
  diag(decomp$sdev[1:14])
Psi hat <- diag(cov(data) - tcrossprod(Lhat))
# Sum squared error
sum((cov(data) - tcrossprod(Lhat) - diag(Psi hat))^2)
## [1] 3.645694
# Compare to the total variance
sum(diag(cov(data)))
```

Example viii

```
## [1] 50.24287

# Our FA model explains:
sum(colSums(Lhat^2)/sum(diag(cov(data))))

## [1] 0.8160488
```

Comments

- In our example above, we saw that 14 factors explained 82% of the total variance.
- ullet Two common default values for m in statistical softwares:
 - Number of positive eigenvalues of the sample covariance matrix.
 - Number of eigenvalues greater than one for the sample correlation matrix.
- In our example, the first criterion would lead to m=p, which is not very helpful

Example (cont'd) i

```
(m <- sum(eigen(cor(data))$values > 1))
## [1] 6
Lhat <- decomp$rotation[,seq_len(m)] %*%
 diag(decomp$sdev[seq_len(m)])
Psi hat <- diag(cov(data) - tcrossprod(Lhat))
# Sum squared error
sum((cov(data) - tcrossprod(Lhat) - diag(Psi hat))^2)
```

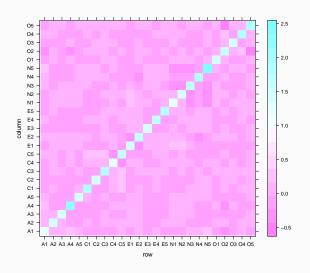
Example (cont'd) ii

```
## [1] 6.999035
# Compare to the total variance
sum(diag(cov(data)))
## [1] 50.24287
# Our FA model explains:
sum(colSums(Lhat^2)/sum(diag(cov(data))))
## [1] 0.5910586
```

Example (cont'd) iii

```
# We can also visualize the fit
Sn <- cov(data)
Sn_fit <- tcrossprod(Lhat) - diag(Psi_hat)
library(lattice)
levelplot(Sn - Sn_fit)</pre>
```

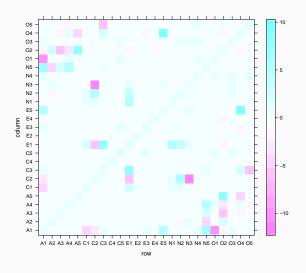
Example (cont'd) iv



Example (cont'd) v

```
# Or if you prefer % difference
levelplot((Sn - Sn_fit)/Sn)
```

Example (cont'd) vi



Estimation-Maximum Likelihood Method i

- This estimation method assumes that both common factors F and specific factors E follow a multivariate normal distribution
 - $\mathbf{F} \sim N_m(0, I)$
 - $\mathbf{E} \sim N_p(0, \Psi)$
- From this assumption, it follows that Y also follows a multivariate normal distribution
 - $\mathbf{Y} \sim N_p(\mu, LL^T + \Psi)$
- Therefore, we can write down the likelihood in terms of both L and Ψ .

Estimation-Maximum Likelihood Method ii

- However, because of the factor rotation problem, we need to impose a constraint in order to obtain a unique solution:
 - $L^T \Psi^{-1} L = \Delta$ is diagonal.
- Given this assumption, the maximum likelihood estimates \hat{L} and $\hat{\Psi}$ can be found using an iterative algorithm.
- We will not go into the details of the algorithm (but see Johnson & Wichern, Supplement 9A if interested).
 - Instead, we will rely on the R function stats::factanal.

Example (cont'd) i

Example (cont'd) ii

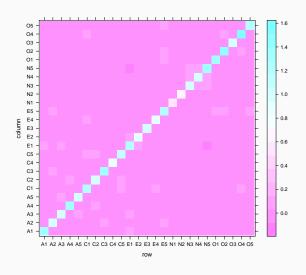
[1] 0.4498342

```
# We get an estimate of the correlation
R mle <- tcrossprod(Lmle) - diag(Psi mle)</pre>
sum((cor(data) - R mle)^2)
## [1] 31.97525
# Our FA model explains:
sum(colSums(Lmle^2)/ncol(data))
```

Example (cont'd) iii

```
levelplot(cor(data) - R_mle)
```

Example (cont'd) iv



Example (cont'd) v

```
# To factor the covariance matrix
# Use psych::fa
fa_decomp <- psych::fa(data, nfactors = m,</pre>
                        rotate = "none",
                        covar = TRUE,
                        fm = "ml")
# Extract estimates
Psi mle <- fa decomp$uniquenesses
Lmle <- fa decomp$loadings
```

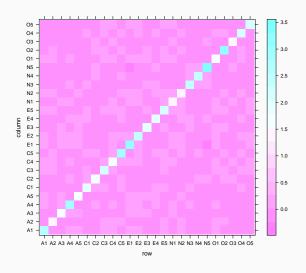
Example (cont'd) vi

```
# We get an estimate of the covariance
Sn mle <- tcrossprod(Lmle) - diag(Psi_mle)</pre>
sum((Sn - Sn mle)^2)
## [1] 128.025
# Our FA model explains:
sum(colSums(Lmle^2)/sum(diag(Sn)))
```

[1] 0.459665

Example (cont'd) vii

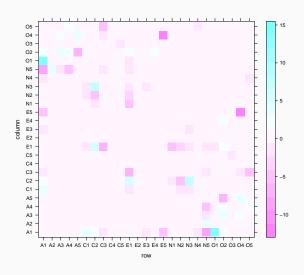
Example (cont'd) viii



Example (cont'd) ix

```
# Compare MLE with PC estimate
levelplot((Sn_fit - Sn_mle)/Sn)
```

Example (cont'd) x



Comments about estimation

- There are other methods of estimating the loadings and the "uniquenesses"
 - Ordinary Least Squares
 - Weighted OLS
 - Principal factor
- With so many choices of estimation methods, it can be hard to compare statistical softwares
 - And you also have to read the manual in order to know what is going on...
- You also always have the choice between factoring the covariance or the correlation matrix
 - Which one you choose depends on the commensurability of your variables (just like in PCA)
- Finally, it's always a good idea to compare the output of

Factor Rotation Redux i

lacktriangleright As we saw earlier, any orthgonal matrix T gives rise to the same factor analysis model

$$\Sigma = LL^T + \Psi = LTT^TL^T + \Psi = \tilde{L}\tilde{L}^T + \Psi.$$

- In other words, we cannot choose T to maximise the goodness of fit.
 - We need another criterion
- Intuitively, to ease interpretation, we want each variable to have large loadings for one factor and negligeable loadings for the other ones.

Factor Rotation Redux ii

- https://maxturgeon.ca/f19stat4690/factor_rotation.gif
- One common analytic criterion that formalises this idea is the varimax criterion.
 - Resulting loadings are called varimax loadings
- We have

$$ext{VARIMAX} \propto \sum_{j=1}^m egin{pmatrix} ext{Variance of squares of scales loadings} \\ ext{for } j\text{-th factor} \end{pmatrix}$$

- More precisely:
 - Let $\tilde{\ell}_{ij}$ be the (i,j)-th entry of the matrix $\tilde{L}=LT$. In other words, $\tilde{\ell}_{ij}$ depends on T.

Factor Rotation Redux iii

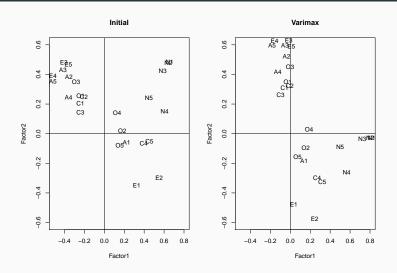
- Let $\tilde{h}_i^2 = \sum_{j=1}^m \tilde{\ell}_{ij}^2$.
- Define the scaled loadings $\tilde{\ell}_{ij}^* = \tilde{\ell}_{ij}/h_i$.
- lacktriangle The varimax criterion V is given as

$$V = \frac{1}{p} \sum_{j=1}^{m} \left(\sum_{i=1}^{p} \tilde{\ell}_{ij}^{*4} - \frac{1}{p} \left(\sum_{i=1}^{p} \tilde{\ell}_{ij}^{*2} \right)^{2} \right).$$

• In R, you can compute the rotated loadings using the stats::varimax function. Alternatively, the function stats::factanal can compute the rotation for you as part of the factor analysis (and so does psych::fa).

Example (cont'd) i

Example (cont'd) ii



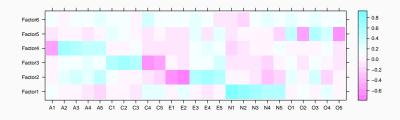
Example (cont'd) iii

[1] 0.6744813

```
# You can extract the matrix T
varimax loadings$rotmat
              \lceil .1 \rceil \qquad \lceil .2 \rceil
##
## [1.] 0.7810310 -0.6244923
## [2,] 0.6244923 0.7810310
# We can also get the angle of rotation
acos(varimax loadings$rotmat[1,1])
```

Example (cont'd) iv

Example (cont'd) v



Comments

- As with estimation, there are many more rotation methods.
 - See for example the help page ?GPArotation::rotations
- One particular class of rotations are called oblique
 - lacktriangle The matrix T is no longer constrained to be orthogonal.
- Factor rotation is especially useful with loadings obtained through MLE
 - Recall the constraint on $L^T \Psi^{-1} L$ being diagonal
- Factor rotations are also sometimes used with PCA loadings.