## **Chapter 4: Factor Analysis**

- In many studies, we may not be able to measure directly the variables of interest.
- We can merely collect data on other variables which may be related to the variables of interest.
- Goal of *factor analysis* (FA) is to relate the unobservable *latent variables* of interest to the observed *manifest variables*.
- The technique used to relate the latent variables (often called *factors*) to the manifest variables is similar to multiple regression.
- The estimation of the regression coefficients (called *loadings* in this context) is less straightforward, however.

## A Factor Analysis Example: The Wechsler Adult Intelligence Study

- The Wechsler Adult Intelligence Scale (WAIS) series of tests measures participants' scores in 11 different tests.
- The multivariate data set consisted of 13 variables: these 11 test scores, plus "age" and "years of education."
- Based on the observed variables, we may want to identify certain underlying factors that cause the individuals to differ.
- Is there a "general intelligence" factor? Is there a "language ability" factor? Is there a "math ability" factor?
- Factor analysis can help us answer these questions.

### **Our Factor Analysis Model**

- Our factor analysis model assumes that we can explain the correlations among the manifest variables through these variables' relationships with the latent variables.
- The q manifest variables are denoted  $x_1, x_2, \ldots, x_q$ .
- The k latent variables, or *factors*, (where k < q) are denoted  $f_1, f_2, \ldots, f_k$ .
- We relate them via a series of regression equations:

$$x_1 = \lambda_{11}f_1 + \lambda_{12}f_2 + \dots + \lambda_{1k}f_k + u_1$$
  

$$x_2 = \lambda_{21}f_1 + \lambda_{22}f_2 + \dots + \lambda_{2k}f_k + u_2$$
  

$$\vdots$$
  

$$x_q = \lambda_{q1}f_1 + \lambda_{q2}f_2 + \dots + \lambda_{qk}f_k + u_q$$

- The  $\lambda_{ij}$  values (called *loadings*) show how much each manifest variable depends on the *j*-th factor.
- The loading values help in the interpretation of each factor.

STAT J530

### **Our Factor Analysis Model (continued)**

• We can write the regression equations in matrix notation:  $\mathbf{x} = \Lambda \mathbf{f} + \mathbf{u}$ , where

$$oldsymbol{\Lambda} = \left[egin{array}{cccc} \lambda_{11} & \cdots & \lambda_{1k} \ dots & \ddots & dots \ \lambda_{q1} & \cdots & \lambda_{qk} \end{array}
ight]$$

and  $\mathbf{f} = (f_1, \dots, f_k)'$ ,  $\mathbf{u} = (u_1, \dots, u_q)'$ .

- The model assumes  $u_1, \ldots, u_q$  are mutually independent and are independent of the  $f_1, \ldots, f_k$ .
- The factors are unobserved, so we may assume they have mean 0 and variance 1, and that they are uncorrelated with each other.

Hitchcock

### Partitioning the Variance of the Data Vectors

The communality  $h_i^2$  is the variability in manifest variable  $x_i$  shared with the other variables (via the factors) and  $\psi_i$  is the specific variance, not shared with the other variables.

Hitchcock

### **Covariance of the Data Vectors**

Hence the population covariance matrix  $\Sigma$  for  $(x_1, x_2, \ldots, x_q)$  is  $\Sigma = \Lambda \Lambda' + \Psi$ , where  $\Psi = diag(\psi_i)$ .

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# **Factor Analysis in Practice**

- If this decomposition of the covariance matrix holds, then the *k*-factor model is correct.
- In practice,  $\Sigma$  is unknown and is estimated by S (or the sample correlation matrix R will be used).
- So we need to find *estimates* of  $\Lambda$  and  $\Psi$  so that the sample covariance matrix can be decomposed in this way:  $\mathbf{S} \approx \hat{\Lambda} \hat{\Lambda}^{'} + \hat{\Psi}$ .
- In practice, we also don't know the true value of k, the number of factors.

# Methods of Estimating the Factor Analysis Model: Principal Factor Analysis

- The *Principal Factor Analysis* approach to estimation relies on estimating the communalities.
- It uses the reduced covariance matrix  $\mathbf{S}^* = \mathbf{S} \hat{\mathbf{\Psi}}$ .
- The diagonal elements of  $S^*$  are  $s_i^2 \hat{\psi}_i = \hat{h}_i^2$ , the (estimated) communality for the *i*-th variable.
- We could standardize the variables, which amounts to using the reduced correlation matrix  ${f R}^*={f R}-\hat{{f \Psi}}.$

## **Estimating the Communalities**

- To estimate the  $h_i^2$  values, we cannot use the factor loadings, since those have not been estimated yet.
- A more straightforward approach (when working with the correlation matrix) is one of the following:
  - 1. Initially let  $\hat{h}_i^2$  equal the  $R^2$  value of a regression of  $x_i$  against the other manifest variables. This is  $1 \frac{1}{r^{ii}}$ , where  $r^{ii}$  is the *i*-th diagonal element of  $\mathbf{R}^{-1}$ .
  - 2. Initially let  $\hat{h}_i^2$  equal the largest absolute correlation coefficient between  $x_i$  and any other manifest variable.
- In both of these approaches, a stronger association between  $x_i$  and the other variables will lead to a higher communality value  $\hat{h}_i^2$ .
- When working with the covariance matrix, we could base the communality estimates on the diagonal elements of  $S^{-1}$  rather than  $R^{-1}$ .

## **Using the Initial Communality Estimates**

- Once we have our initial  $\hat{h}_i^2$  values, we can calculate  $\mathbf{S}^*$  (or  $\mathbf{R}^*$ ).
- We perform a principal components analysis on  $S^*$  (or  $R^*$ ) and the first k eigenvectors contain the estimates of the first k factor loadings.
- These estimated loadings  $\hat{\lambda}_{ij}$  can be used to obtain new communality estimates:

$$\hat{h}_i^2 = \sum_{j=1}^k \hat{\lambda}_{ij}^2$$

- We can re-form  $S^*$  (or  $R^*$ ) with the revised communality estimates, and repeat the process until the communality estimates converge.
- This works well unless the communality estimate becomes larger than the manifest variable's total variance, implying a negative specific variance, an impossibility.

## Maximum Likelihood Factor Analysis

- Maximum likelihood (ML) is a general method of estimating parameters in a statistical model.
- Classical ML requires an assumption about the form of the distribution of the data.
- If we can assume we have multivariate normal data, we can motivate a maximum likelihood estimation of our k-factor model.
- Recall that the observed sample covariance matrix is  ${f S}$  and, under the factor analysis model, the true covariance matrix is  ${f \Sigma}=\Lambda\Lambda^{'}+{f \Psi}.$
- The goodness-of-fit of the *k*-factor model can be judged by a "distance" measure *F* between the sample covariance matrix and the predicted covariance matrix under the model.

#### The Distance Measure and Maximum Likelihood

- Let  $F = \ln |\Lambda \Lambda' + \Psi| + trace(\mathbf{S}[\Lambda \Lambda' + \Psi]^{-1}) \ln |\mathbf{S}| q.$
- This distance measure equals zero if  $\mathbf{S}=\mathbf{\Lambda}\mathbf{\Lambda}^{'}+\mathbf{\Psi}.$
- F is large when  ${f S}$  is far from  ${f \Lambda \Lambda}' + {f \Psi}.$
- We can calculate (for a given data set) the elements of  $\Lambda$  and  $\Psi$  that make F as small as possible.
- This implies we have estimates of the communalities  $h_1^2, \ldots, h_q^2$  and the specific variances  $\psi_1, \ldots, \psi_q$ .
- Under the assumption of multivariate normality, the likelihood L = -0.5 nF plus a function of the data.
- Hence minimizing F is equivalent to maximizing L.
- This method could also produce negative estimates for the specific variances.

## **Estimating the Number of Factors**

- With factor analysis, the choice of the number of factors k is critical.
- If we use k + 1 factors, we will get different factors and loadings than if we use k factors.
- With too few factors, there will be too many high loadings.
- With too many factors, the loadings will be spread out too much over the factors, and the factors will be difficult to interpret.

### Methods for Estimating the Number of Factors

- A subjective approach is to try various choices of k and pick the one that gives the most interpretable result this is probably *too* subjective.
- Could use the *scree diagram* as in PCA, but the eigenvalues are not as directly interpretable in factor analysis.
- When using maximum likelihood, we can use a formal sequence of hypothesis tests to help determine *k*.
- We use the test statistic  $U = n' \min(F)$ , where n' = n + 1 (2q+5)/6 2k/3.
- If the k-factor model is appropriate, this test statistic has a large-sample  $\chi^2$  distribution with degrees of freedom  $(q k)^2/2 (q + k)/2$ .
- Typically we begin with a small value of k, and increase k by 1 sequentially.
- If at any stage, the U has a non-significant P-value, we choose that value of k.
- If at any stage the degrees of freedom go to zero, the factor analysis model may be inappropriate.