Principal Component Analysis Factor Analysis



Bird's-eye (over)view of existing mixtures methods



Outline

- Introduction
- PCA
- 3 FA
- 4 Notes

Introduction

- While clustering aims to identify distinct subgroups based on exposure experience
- Factor analytic techniques aim to describe variability among observed, correlated chemicals in terms of a potentially lower number of unobserved factors
- These two unsupervised approaches are actually connected, but we will not get into this here
 - If interested in this please read: Ding C, He X. "K-means clustering via principal component analysis." ICLM 2004

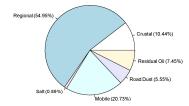
Introduction PCA FA Note

Pattern Recognition & Dimensionality Reduction

- A potential topic of interest in assessing exposure to mixtures is to identify patterns of exposure
- This can be either specific sources of exposure
- Or common behaviors in the study population
- These patterns, to be meaningful, tend to have a *lower dimension* than the original data matrix, i.e. k < p
- The most common example arises in air pollution
 - Source apportionment

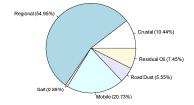
Source Apportionment

- \circ Use factor analytic techniques to apportion $\text{PM}_{2.5}$ to its sources
 - E.g. Boston 2003 2010
 - $\circ~$ BC and 17 other $PM_{2.5}$ XRF-measured components



Source Apportionment

- Use factor analytic techniques to apportion PM_{2.5} to its sources
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- Use PM_{2.5} sources as the exposure of interest in the health models
 - Identification of most harmful emission sources
 - More effective air quality management and regulations

Pattern Recognition & Dimensionality Reduction (cont'd)

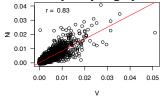
- There are many approaches for pattern recognition and dimensionality reduction
- Most common applications are in Computer Science and Machine Learning (e.g. face recognition etc)
- And many are being developed
- Here we will focus on the two most "traditional"
 - Principal Component Analysis (PCA)
 - Pactor Analysis (FA)

What is PCA?

- Dimensionality reduction statistical tool
- Aims to explain as much of the total variance in the data as possible using a smaller number of variables
- The resulting variables (i.e. components) are linear combinations of the original variables
- Reducing the dimension of the data will somewhat reduce accuracy – tradeoff

2-D Example

- Ni and V are associated with residual oil combustion
- Commonly very highly correlated



- The above points are presented in two axes
- PCA will allow us to present the data along one axis (the red in this case), called *principal component*
- o In reality, we don't use PCA for 2-D problems

How does PCA Work?

PCA aims to:

- Identify a sequence of linear combinations of the p variables
- That have maximal variance
- And are mutually uncorrelated
 - I.e. orthogonal solution

Eigenvalues & Eigenvectors

$$A\nu = \lambda \nu$$

where A is the $p \times p$ var-covar or correlation matrix of X, ν is the eigenvector $(p \times 1)$ and λ is a scalar (eigenvalue)

- Every eigenvector has an eigenvalue (p pairs)
- The eigenvector shows the direction (i.e. in the 2-D example above the direction of the red line)
- The eigenvalue is a scalar saying how much variance exists in the data along that direction
- The eigenvector with the highest eigenvalue is the first principal component

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Singular Value Decomposition (SVD)

- SVD is a generalization of the eigen decomposition that also works on non-square matrices
 - I.e. on the data matrix directly
- Preferable to standardize (center and scale) variables
 - Especially if chemicals are on different scales and/or units
 - Most software has options to automatically do this
- Both SVD and eigen decomposition yield orthogonal solutions
 - I.e. the estimated components are not correlated

PCA – Results

$$z_{i1} = \phi_{11}x_{i1} + \phi_{21}x_{i2} + \dots + \phi_{p1}x_{ip}$$

$$z_{i2} = \phi_{12}x_{i1} + \phi_{22}x_{i2} + \dots + \phi_{p2}x_{ip}$$

$$\vdots$$

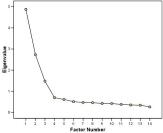
$$z_{ip} = \phi_{1p}x_{i1} + \phi_{2p}x_{i2} + \dots + \phi_{pp}x_{ip}$$

- \circ $i \in 1, \cdots, N$
- \circ z_{11}, \cdots, z_{N1} the scores on the first PC
- $\phi_1 = (\phi_{11}, \phi_{21}, \cdots, \phi_{p1})^T \rightarrow \text{loadings of the first PC}$
- $\phi_1, \cdots, \phi_p \to \text{eigenvectors}$

$$\circ \sum_{j=1}^{p} \phi_{j1}^2 = 1$$

PCA – Dimensionality Reduction

- PCA will yield a new data matrix of the same dimensions $(N \times p)$
- We need to choose how many components to keep for further analysis
- Look for "elbow" at scree plot



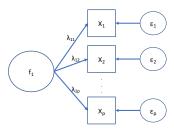
 Decide how much of the total variance explained we'd like to keep

Factor Analysis (FA)

- FA assumes that the measurements of the p pollutants arise from k underlying sources
 - That are not observed (i.e. latent)
 - With k < p, and k pre-specified
- Does not require orthogonality
- What we will discuss here is exploratory FA (EFA)
 - SEMs lie under confirmatory FA (CFA)

FA (cont'd)

- $\mathbf{x} = \boldsymbol{\mu} + \boldsymbol{\Lambda} \boldsymbol{f} + \boldsymbol{\varepsilon}$, where
 - Λ a $p \times k$ matrix of factor loadings
 - f a matrix of k common sources of variation among X and accounts for their correlation structure
 - \circ arepsilon have mean zero and Ψ diagonal covariance matrix



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FA (cont'd)

- To solve FA we can
- Assume joint normality and use maximum likelihood
 - Ends up being a hard optimization problem
 - With often many local minima
- SVD of the data matrix
 - o If we assume that instead of diagonal Ψ the error matrix is Ξ and small (in sums of squares of its elements)
 - \circ Now just a question of minimizing sums of squares \rightarrow reduces to matrix decompositions
 - Rotation to yield more interpretable factors (potentially correlated)
 - If the factor correlations are not driven by the data the solution will remain nearly orthogonal

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- PCA aims to explain the total variance in the data
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- In PCA all p components are provided and the user selects k afterwards
- \circ In FA k needs to be pre-specified
- Usually similar solutions

- For both PCA & FA: no single correct answer
 - \circ No "golden rule" to select k
- $\circ\,$ And scores and factors are centered at zero \cdots
- Interpretability is key (and an issue)

troduction PCA FA **Notes**

PCA and FA in Health Models

- When the components/factors have been identified the resulting scores or source contributions can be included as the exposure(s) of interest in health models
- The PCA solution is orthogonal no need to worry about co-component confounding
- This might not be the case with FA
- \circ These are now continuous \to non-linear exposure-response functions can also be explored
- Uncertainty propagation?
- Supervised extensions exist

Thank you!

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