Model-Based Clustering III

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1

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Introduction

- Now, we are going to look at clustering longitudinal data in a mixture framework.
- As in the last couple of lectures, some of the material is taken from McNicholas (2016).
- Bibliographic references are given at the end of these slides.

Cholesky Decomposition

- The Cholesky decomposition is a method for decomposing a matrix into the product of a lower triangular matrix and its transpose.
- Let A be a real, positive definite matrix; then the Cholesky decomposition of A is

$$\mathbf{A} = \mathbf{L}\mathbf{L}',\tag{1}$$

where L is a unique lower triangular matrix.

• The decomposition in (1) is often used in numerical analysis applications to simplify the solution to a system of linear equations.

3

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Modified Cholesky Decomposition

• Pourahmadi (1999) applies a modified Cholesky decomposition to the covariance matrix Σ of a random variable, i.e.,

$$\mathbf{T}\mathbf{\Sigma}\mathbf{T}' = \mathbf{D},\tag{2}$$

where T is a unique unit lower triangular matrix and D is a unique diagonal matrix with strictly positive diagonal entries.

- Note that a unit lower triangular matrix is a lower triangular matrix with diagonal elements 1.
- \bullet Alternatively, the relationship in (2) can be written

$$\mathbf{\Sigma}^{-1} = \mathbf{T}' \mathbf{D}^{-1} \mathbf{T}. \tag{3}$$

Modified Cholesky Decomp. contd.

• The values of T and D can be interpreted as generalized autoregressive parameters and innovation variances, respectively (Pourahmadi, 1999), so that the linear least-squares predictor of X_t , based on X_{t-1}, \ldots, X_1 , is given by

$$\hat{X}_t = \mu_t + \sum_{s=1}^{t-1} (-\varphi_{ts})(X_s - \mu_s) + \sqrt{d_t}\varepsilon_t, \tag{4}$$

where φ_{ts} is the sub-diagonal element of \mathbf{T} in position (t,s), d_t is the tth diagonal element of \mathbf{D} , and $\varepsilon_t \sim \mathsf{N}(0,1)$.

 To cluster longitudinal data, McNicholas and Murphy (2010) use (3) as the component covariance in a Gaussian mixture, i.e.,

$$\mathbf{\Sigma}_g^{-1} = \mathbf{T}_g' \mathbf{D}_g^{-1} \mathbf{T}_g.$$

5

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A Family

- A family of eight Gaussian mixture models arises from the option to constrain \mathbf{T}_g and/or \mathbf{D}_g to be equal across components together with the option to impose the isotropic constraint $\mathbf{D}_g = \delta_g \mathbf{I}_p$ (McNicholas and Murphy, 2010).
- Each member of this family has a natural interpretation for longitudinal data
 - Constraining $T_g = T$ suggests that the autoregressive relationship between time points, cf. (4), is the same across components.
 - The constraint $\mathbf{D}_g = \mathbf{D}$ suggests that the variability at each time point is the same for each component.
 - The isotropic constraint $\mathbf{D}_g = \delta_g \mathbf{I}_p$ suggests that the variability is the same at each time point in component g.

6

A Family contd.

Model	\mathbf{T}_g	\mathbf{D}_g	\mathbf{D}_g	Free Cov. Parameters
EEA	Equal	Equal	Anisotropic	p(p-1)/2 + p
VVA	Variable	Variable	Anisotropic	G[p(p-1)/2] + Gp
VEA	Variable	Equal	Anisotropic	G[p(p-1)/2] + p
EVA	Equal	Variable	Anisotropic	p(p-1)/2 + Gp
VVI	Variable	Variable	Isotropic	G[p(p-1)/2] + G
VEI	Variable	Equal	Isotropic	G[p(p-1)/2] + 1
EVI	Equal	Variable	Isotropic	p(p-1)/2 + G
EEI	Equal	Equal	Isotropic	p(p-1)/2 + 1

7

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Comments

- Note that an EM algorithm can be used for parameter estimation.
- I want to look at some clustering examples now.
- We can use the longclust package in R.
- Here, it is not so easy to compare with other approaches.
- Among other things, I hope you are getting a sense of the flexibility of mixture model-based approaches.
- In the next lecture, we will finish up with model-based approaches and move on.

8

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References

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- Pourahmadi, M. (1999). 'Joint mean-covariance models with applications to longitudinal data: Unconstrained parameterisation'. *Biometrika* 86(3), 677–690.

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