

Final exam: Tuesday April 19 7-9 pm
 Coverage: Chap. 1-6 of text + classification
 - 2 (5.5"x11") aid sheets (both sides) and a calculator are allowed.

Office hrs: Next week TBA
 Mon April 18 1-3 pm

Structured multivariate Models

So far: $\underline{X} \sim N_p(\underline{\mu}, C)$

Graphical model: $K = C^{-1} \rightarrow$ assume that K is sparse — many 0 entries

PCA: C can be approximated by a lower rank matrix.

Factor Analysis: $C = L L^T + \Psi \rightarrow$ diagonal
 $\downarrow p \times r$

More generally: Apply multivariate thinking to univariate models

Example: Longitudinal / panel data

n subjects followed over time

— measure some (univariate) response at time pts t_1, \dots, t_p

X_{ij} = response for subject i at time t_j
 $= f(t_j) + \text{error}$

$\underline{X}_i = \begin{pmatrix} X_{i1} \\ \vdots \\ X_{ip} \end{pmatrix} \rightarrow$ model dependence of observations within each subjects.

Example: Paired difference design

- 2 treatments
- n subjects

Each subject receives both treatments
 \swarrow subject effect (random)

Model: $X_{ij} = \mu_i + \mu_j + \varepsilon_{ij} \rightarrow$ noise
 $\swarrow \quad \searrow \quad \downarrow$
 subject treatment treatment effect (fixed)

Look at $\Delta_i = X_{i1} - X_{i2} = \underbrace{\mu_1 - \mu_2}_{\text{treatment difference}} + \underbrace{\varepsilon_{i1} - \varepsilon_{i2}}_{\text{noise}}$

For example, the test $H_0: \mu_1 = \mu_2$ or obtain a CI for $\mu_1 - \mu_2$, we use $\Delta_1, \dots, \Delta_n$

Back to original data

$$\underline{X}_i = \begin{pmatrix} X_{i1} \\ X_{i2} \end{pmatrix} = \mu_i \begin{pmatrix} 1 \\ 1 \end{pmatrix} + \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix} + \begin{pmatrix} \varepsilon_{i1} \\ \varepsilon_{i2} \end{pmatrix}$$

Assume $A_i \sim N(0, \sigma_A^2)$ (independent)

$\underline{\varepsilon}_i \sim N_2(0, \sigma_\varepsilon^2 I)$ (independent)

A_i independent of $\underline{\varepsilon}_i$

$\Rightarrow \underline{X}_i \sim N_2(\underline{\mu}, C)$ where $C = \text{Cov}(A_i(1), \underline{\varepsilon}_i) = \text{Cov}(A_i(1)) + \text{Cov}(\underline{\varepsilon}_i)$

$$C = \sigma_A^2 \begin{pmatrix} 1 \\ 1 \end{pmatrix} \begin{pmatrix} 1 & 1 \end{pmatrix} + \sigma_\varepsilon^2 \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}$$

$$= \begin{pmatrix} \sigma_A^2 + \sigma_\varepsilon^2 & \sigma_A^2 \\ \sigma_A^2 & \sigma_A^2 + \sigma_\varepsilon^2 \end{pmatrix}$$

$$\text{Corr}(X_{i1}, X_{i2}) = \frac{\sigma_A^2}{\sigma_A^2 + \sigma_\varepsilon^2}$$

Note: C has a single factor form $C = \underline{L} \underline{L}^T + \Psi$

A step further: Repeated measures design

$\left. \begin{array}{l} k \text{ treatments} \\ n \text{ subjects} \end{array} \right\}$ each subject receives all k treatments

— "treatment" used loosely here — for example, we could have a response measured at different points in time.

$$\underline{X}_i = \begin{pmatrix} X_{i1} \\ \vdots \\ X_{ik} \end{pmatrix} = A_i \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} + \begin{pmatrix} \mu_1 \\ \vdots \\ \mu_k \end{pmatrix} + \begin{pmatrix} \varepsilon_{i1} \\ \vdots \\ \varepsilon_{ik} \end{pmatrix}$$

$$= A_i \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} + \underline{\mu} + \underline{\varepsilon}_i$$

Assumption: $A_i \sim N(0, \sigma_A^2)$ (independent)

$\underline{\varepsilon}_i \sim N_k(0, \sigma_\varepsilon^2 I)$ (indep)

$A_i + \underline{\varepsilon}_i$ independent

$$\underline{X}_i \sim N(\underline{\mu}, C) \text{ where } C = \begin{pmatrix} \sigma_A^2 + \sigma_\varepsilon^2 & \sigma_A^2 & \cdots & \sigma_A^2 \\ \vdots & \ddots & \ddots & \vdots \\ \sigma_A^2 & \cdots & \cdots & \sigma_A^2 + \sigma_\varepsilon^2 \end{pmatrix} \leftarrow C \text{ is defined by 2 parameters } \sigma_A^2, \sigma_\varepsilon^2$$

Estimation: $\mu_1, \dots, \mu_k, \sigma_\varepsilon^2, \sigma_A^2$

① Likelihood estimation of all parameters

— full likelihood

$$L(\mu_1, \dots, \mu_k, \sigma_A^2, \sigma_\varepsilon^2) = \prod_{i=1}^n \left\{ \frac{1}{(2\pi)^{k/2} |C|^{1/2}} \exp \left[-\frac{1}{2} (\underline{X}_i - \underline{\mu})^T C^{-1} (\underline{X}_i - \underline{\mu}) \right] \right\}$$

dependent on $\sigma_A^2, \sigma_\varepsilon^2$

Restricted ML estimation (REML)

— transform data to eliminate $\mu_1, \dots, \mu_k \rightarrow$ get estimate of $\sigma_A^2, \sigma_\varepsilon^2$

— Then estimate μ_1, \dots, μ_k by generalized least squares.

② Transform data to eliminate random effects A_1, \dots, A_n

$$\underline{X}_i = A_i \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} + \underline{\mu} + \underline{\varepsilon}_i$$

Multiply by some $(k-1) \times k$ matrix B such that $B \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} = \underline{0}$

$$\underbrace{BX_i}_{X_i^*} = \underbrace{B\mu}_{\mu^*} + \underbrace{B\varepsilon_i}_{\varepsilon_i^*} \quad N_{k+1}(Q, \sigma_\varepsilon^2 BB^T)$$

Examples

$$B = \begin{pmatrix} 1 & -1 & 0 & \dots & 0 \\ 0 & 1 & -1 & & 0 \\ 0 & 0 & 1 & -1 & \\ 0 & \dots & & 1 & -1 \end{pmatrix}$$

$$B\mu = \begin{pmatrix} \mu_1 - \mu_2 \\ \mu_2 - \mu_3 \\ \vdots \\ \mu_{k-1} - \mu_k \end{pmatrix}$$

$$B = \begin{pmatrix} -1 & 0 & 0 & \dots & 0 \\ -1 & 0 & 1 & & 0 \\ \vdots & & & \ddots & \\ -1 & 0 & \dots & 1 \end{pmatrix}$$

$$B\mu = \begin{pmatrix} \mu_2 - \mu_1 \\ \vdots \\ \mu_k - \mu_1 \end{pmatrix}$$

Advantage: Eliminate random effect

Disadvantage: Lose information in transformation

→ can't estimate μ itself but B instead.