

Lecture 3 | 11.03.2024

# Statistical inference in a multivariate regression model

# Notation overview

- balanced longitudinal profiles  $\mathcal{D}_B \equiv \{(\mathbf{Y}_i, \mathbf{X}_{i1}^\top, \dots, \mathbf{X}_{in}^\top)^\top; i = 1, \dots, N\}$ 
  - for  $n_i = n \in \mathbb{N}$  for all  $i = 1, \dots, N$
  - random vectors  $(\mathbf{Y}_i, \mathbf{X}_{i1}^\top, \dots, \mathbf{X}_{in}^\top)^\top$  are independent with the same length
  - for **longitudinal data** we do not assume that subject specific measurements are taken at the same time  $\Rightarrow \mathcal{D}_B$  **generally not a random sample!**
  - for **multivariate regression model** we already assume that the observations in  $\mathcal{D}_B$  **form a random sample**  $\Rightarrow$  notation  $\mathcal{D}_S$
- population and data model formulation (theoretical vs. empirical)

$$\mathbf{Y} = \mathbf{X}^\top \mathbb{B} + \boldsymbol{\varepsilon} \qquad \mathbb{Y} = \mathbb{X} \mathbb{B} + \mathbb{U}$$

for generic random vectors  $\mathbf{Y} \in \mathbb{R}^n$  and  $\mathbf{X} \in \mathbb{R}^p$  and some matrix of the **unknown parameters**  $\mathbb{B} \in \mathbb{R}^{p \times n}$

The corresponding data:  $\mathbb{Y} = (\mathbf{Y}_1^\top, \dots, \mathbf{Y}_N^\top)^\top$ ,  $\mathbb{X} = (\mathbf{X}_1^\top, \dots, \mathbf{X}_N^\top)^\top$ , and  $\mathbb{U} = (\mathbf{u}_1^\top, \dots, \mathbf{u}_N^\top)^\top \equiv (\boldsymbol{\varepsilon}_1^\top, \dots, \boldsymbol{\varepsilon}_N^\top)^\top$

## Statistical inference: Likelihood ratio test

- Inference in terms of **confidence intervals/regions** and **hypothesis tests**
- General form of the null hypothesis:

$$H_0 : \mathbb{C}_1 \mathbb{B} \mathbb{M}_1 = \mathbb{D}$$

where  $\mathbb{C}_1$ ,  $\mathbb{M}_1$ , and  $\mathbb{D}$  are some (suitable) matrices

- The rows of  $\mathbb{C}_1$  do inference on the effects of independent variables while the columns of  $\mathbb{M}_1$  do inference on particular linear combinations of dependent variables
- In practical applications it is common that  $\mathbb{D}$  is a zero matrix (all elements are zeros) and  $\mathbb{M}_1 = \mathbb{I}$  (i.e. a unit matrix with ones on the main diagonal)   
 $\hookrightarrow$  alternatively, the model of the form  $\mathbb{Y} \mathbb{M}_1 = \mathbb{X} \mathbb{B} \mathbb{M}_1 + \mathbb{U} \mathbb{M}_1$
- Thus, the null hypothesis reduces to

$$H_0 : \mathbb{C}_1 \mathbb{B} = \mathbf{0}$$

against a general alternative hypothesis of the form  $H_A : \mathbb{C}_1 \mathbb{B} \neq \mathbf{0} \in \mathbb{R}^{q \times n}$  (with the rank of the matrix  $\mathbb{C}_1$  being equal to  $q \in \mathbb{N}$ )

## Inference: Likelihood ratio test

- consider the null hypothesis of the form  $H_0 : C_1 B = D$
- the model  $Y = XB + U$  can be equivalently expressed as

$$\tilde{Y} = Z\tilde{B} + U,$$

for  $\tilde{Y} = Y - XB_0$ , where  $C_1 B_0 = D$  (satisfies the null hypothesis),  
 $Z = XC^{-1}$  where  $C^T = (C_1^T, C_2^T)$  and  $\tilde{B} = (\tilde{B}_1^T, \tilde{B}_2^T)^T = C(B - B_0)$

- the null hypothesis  $C_1 B = D$  gives that  $\tilde{B}_1 = \mathbf{0}$  and for the matrix partition  $C^{-1} = (C^{(1)}, C^{(2)})$  the projection matrix

$$P_1 = I - XC^{(2)}(C^{(2)T}X^TXC^{(2)})^{-1}C^{(2)T}X^T$$

defines the projection onto the linear subspace orthogonal to the columns of the matrix  $XC^{(2)}$  (i.e., residuals for the regression onto  $C^{(2)}$  – under the null hypothesis, thus  $\tilde{B}_1 = \mathbf{0}$ )

# LRT: Likelihood under the null and alternative

- maximized likelihood under the null hypothesis

$$\ell_0 = |2\pi N^{-1} \tilde{Y}^\top P_1 \tilde{Y}|^{-N/2} \cdot \exp\left\{-\frac{1}{2} Nn\right\}$$

- maximized likelihood under the alternative hypothesis

$$\ell_1 = |2\pi N^{-1} \tilde{Y}^\top \tilde{P} \tilde{Y}|^{-N/2} \cdot \exp\left\{-\frac{1}{2} Nn\right\}$$

- the likelihood ratio test statistic is given as

$$\lambda^{2/N} = |\tilde{Y}^\top \tilde{P} \tilde{Y}| / |\tilde{Y}^\top P_1 \tilde{Y}| = |\tilde{Y}^\top \tilde{P} \tilde{Y}| / |\tilde{Y}^\top \tilde{P} \tilde{Y} + \tilde{Y}^\top P_2 \tilde{Y}|$$

and it follows the  $\Lambda(n, N - p, q)$  distribution, where  $q \in \mathbb{N}$  is the number of rows in  $\mathbb{C}_1$  (for  $P_2 = P_1 - \tilde{P}$  - **what does it mean geometrically?**)

# Examples

- Repeated measurements for two groups (two-sample problems):

$$\mathbf{Y}_i^{(1)} \sim N_n(\boldsymbol{\mu}_1, \boldsymbol{\Sigma}), \quad i = 1, \dots, N_1$$

$$\mathbf{Y}_i^{(2)} \sim N_n(\boldsymbol{\mu}_2, \boldsymbol{\Sigma}), \quad i = 1, \dots, N_2$$

- Typical testing problems:

- parallel profiles of two groups
- identical profiles for both groups
- treatment effect

$$H_0 : \mathbb{C}(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2) = \mathbf{0}$$

$$H_0 : \mathbf{1}^\top (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2) = 0$$

$$H_0 : \mathbb{C}(\boldsymbol{\mu}_1 + \boldsymbol{\mu}_2) = \mathbf{0}$$

- Multiple testing problem:** testing for identical profiles only makes sense if the profiles are parallel; Similarly, if the profiles are parallel, is there any treatment effect at all?

## Two sample problems:

### □ Parallel profiles

$$T = \frac{N_1 N_2}{(N_1 + N_2)^2} (N_1 + N_2 - 2) \left[ \mathbf{C}(\bar{Y}^{(1)} - \bar{Y}^{(2)}) \right]^\top \left( \mathbf{CSC}^\top \right)^{-1} \left[ \mathbf{C}(\bar{Y}^{(1)} - \bar{Y}^{(2)}) \right]$$

and (under the null hypothesis)  $T \sim T^2(n-1, N_1 + N_2 - 2)$

### □ Equality of two levels

$$T = \frac{N_1 N_2}{(N_1 + N_2)^2} (N_1 + N_2 - 2) \frac{\left[ \mathbf{1}^\top (\bar{Y}^{(1)} - \bar{Y}^{(2)}) \right]^2}{\mathbf{1}^\top \mathbf{S1}}$$

and (under the null hypothesis)  $T \sim T^2(1, N_1 + N_2 - 2)$

### □ Same treatment effect

$$T = (N_1 + N_2 - 2) (\mathbf{C}\bar{Y})^\top \left( \mathbf{CSC}^\top \right)^{-1} \mathbf{C}\bar{Y}, \quad \text{for } \bar{Y} = \frac{N_1 \bar{Y}^{(1)} + N_2 \bar{Y}^{(2)}}{N_1 + N_2}$$

and (under the null hypothesis)  $T \sim T^2(n-1, N_1 + N_2 - 2)$

# Overview

- statistical test about some (multivariate) mean vector  $\boldsymbol{\mu} \in \mathbb{R}^n$  can be often expressed in terms of the null hypothesis  $H_0: \mathbb{A}\boldsymbol{\mu} = \mathbf{a}$  vs.  $H_A: \mathbb{A}\boldsymbol{\mu} \neq \mathbf{a}$ , where  $\mathbb{A} \in \mathbb{R}^{q \times n}$  and  $\mathbf{a} \in \mathbb{R}^q$
- for  $\mathbf{X}_i \sim N_n(\boldsymbol{\mu}, \Sigma)$  for  $i = 1, \dots, N$ , with  $\Sigma$  known, the log-likelihood based test statistic  $-2 \log \lambda = N(\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a})^\top (\mathbb{A}\Sigma\mathbb{A}^\top)^{-1}(\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a})$  follows (exactly) the  $\chi^2$  distribution with  $q \in \mathbb{N}$  degrees of freedom
- for  $\mathbf{X}_i \sim N_n(\boldsymbol{\mu}, \Sigma)$  for  $i = 1, \dots, N$ , with  $\Sigma$  unknown, the log-likelihood test statistic  $-2 \log \lambda = N \log \left\{ 1 + (\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a})^\top (\mathbb{A}\hat{\Sigma}_N\mathbb{A}^\top)^{-1}(\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a}) \right\}$  follows asymptotically the  $\chi^2$  distribution with  $q \in \mathbb{N}$  degrees of freedom and the exact Hotelling test is based on the test statistic

$$(N-1)(\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a})^\top (\mathbb{A}\hat{\Sigma}_N\mathbb{A}^\top)^{-1}(\mathbb{A}\bar{\mathbf{X}}_N - \mathbf{a}) \sim T^2(q, N-1)$$

- for  $Y_i \sim N(\mathbf{X}_i^\top \boldsymbol{\beta}, \sigma^2)$ , for  $i = 1, \dots, N$ , with  $\sigma^2 > 0$  unknown, the test of the null hypothesis  $H_0: \mathbb{A}\boldsymbol{\beta} = \mathbf{a}$ , for  $\mathbf{a} \in \mathbb{R}^q$ , leads to the test statistic

$$\frac{N-n}{q} \cdot \frac{(\hat{\boldsymbol{\beta}} - \mathbf{a})^\top \left[ \mathbb{A}(\mathbf{X}^\top \mathbf{X})^{-1} \mathbb{A} \right]^{-1} (\hat{\boldsymbol{\beta}} - \mathbf{a})}{(\mathbf{Y} - \mathbf{X}\hat{\boldsymbol{\beta}})^\top (\mathbf{Y} - \mathbf{X}\hat{\boldsymbol{\beta}})} \sim F_{q, N-n}$$



# Multivariate model vs. general linear model

- ❑ **Multivariate regression model  $\mathbf{Y} = \mathbf{X}\mathbf{B} + \mathbf{U}$** 
  - ❑  $n \in \mathbb{N}$  repeated measurements within  $N \in \mathbb{N}$  subjects (random sample)
  - ❑ repeated measurements taken at the same time-points across subjects
  - ❑ time evolution modeled by the set of  $\beta_j \in \mathbb{R}^P$  parameters ( $j = 1, \dots, n$ )
  - ❑ the vector of subject's specific covariates  $\mathbf{X}_i \in \mathbb{R}^P$  fixed over time
  - ❑ covariance structure modeled by the matrix  $\Sigma$ , where  $\mathbf{u}_i \sim N_n(\mathbf{0}, \Sigma)$
  - ❑ the data usually form a random sample from the joint distribution  $F_{\mathbf{Y}, \mathbf{X}}$
  
- ❑ **General linear model for correlated errors  $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$** 
  - ❑  $n \in \mathbb{N}$  repeated measurements within  $N \in \mathbb{N}$  subjects (balanced data)
  - ❑ the vector of unknown parameters  $\boldsymbol{\beta} \in \mathbb{R}^P$  is fixed over time
  - ❑ subject's specific covariates  $\mathbf{X}_{ij} \in \mathbb{R}^P$  may vary with  $j \in \{1, \dots, n\}$
  - ❑ subjects' independence and within subject's covariance modeled by the variance covariance  $\Sigma$ , where  $\boldsymbol{\varepsilon} \sim N(\mathbf{0}, \Sigma)$  (overall dimensionality:  $Nn$ )
  - ❑ the model can be further generalized for unbalanced data ( $n_i \in \mathbb{N}$ )

## General linear model with correlated errors

- instead of time-varying  $\beta_j$  and fixed  $\mathbf{X}_j \in \mathbb{R}^p$  the time evolution can be modeled in terms of time-varying covariates  $\mathbf{X}_{ij} \in \mathbb{R}^p$  and fixed  $\beta \in \mathbb{R}^p$
- Simplification in terms of the vectors of unknown parameters  $\beta_j \in \mathbb{R}^p$  for  $j = 1, \dots, n$  (in the matrix  $\mathbb{B} \in \mathbb{R}^{p \times n}$ ):  $\Rightarrow \beta = \beta_1 = \dots = \beta_n$
- Relaxation in terms of the subject's specific covariates  $\mathbf{X}_{ij} \in \mathbb{R}^p$  that are now allowed to change with  $j \in \{1, \dots, n\}$ :  $\Rightarrow \mathbf{X}_{ij} = (X_{ij1}, \dots, X_{ijp})^\top \in \mathbb{R}^p$
- this allows for an alternative formulation of the multivariate (data) model (where  $\mathbf{Y} = \mathbf{X}\mathbb{B} + \mathbf{U}$  follows as a special case) in a form

$$\begin{pmatrix} Y_{11} \\ \vdots \\ Y_{1n} \\ Y_{21} \\ \vdots \\ Y_{Nn} \end{pmatrix} = \begin{pmatrix} X_{111} & \dots & X_{11p} \\ \vdots & \ddots & \vdots \\ X_{1n1} & \dots & X_{1np} \\ X_{211} & \dots & X_{21p} \\ \vdots & \ddots & \vdots \\ X_{Nn1} & \dots & X_{Nnp} \end{pmatrix} \begin{pmatrix} \beta_1 \\ \vdots \\ \beta_p \end{pmatrix} + \begin{pmatrix} \varepsilon_{11} \\ \vdots \\ \varepsilon_{1n} \\ \varepsilon_{21} \\ \vdots \\ \varepsilon_{Nn} \end{pmatrix}$$

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- What are the advantages and disadvantages of both model formulations?

# Matrix formulation

- typically we use the notation (under multivariate normal assumption)

$$\mathbf{Y} \sim N_{Nn}(\mathbb{X}\boldsymbol{\beta}, \sigma^2\mathbb{V}),$$

where  $\mathbb{V}$  is a block-diagonal matrix with non-zero blocks of size  $n \times n$  (each block  $\sigma^2\mathbb{V}_0$  represents the variance-covariance of repeated measurements within a single subject)

- the variance covariance matrix  $\sigma^2\mathbb{V}$  is estimated by borrowing power across subject (i.e., replication of  $\sigma^2\mathbb{V}_0$  across the units)
- there can be no specific (parametric) structure assumed for  $\mathbb{V}_0$  but it is common to postulate some parametric form of  $\mathbb{V}_0$
- the correlation structure within  $\sigma^2\mathbb{V}$  is crucial for a proper inference

# Uniform correlation model

- **Assumption:** positive correlation  $\rho \in (0, 1)$  between any two repeated observations within a given subject
- **Matrix notation:**  $\mathbb{V}_0 = (1 - \rho)\mathbb{I}_{n \times n} + \rho\mathbf{1}_{n \times n}$
- **Motivation:** the response (random) variable  $Y_{ij}$  can be decomposed as

$$Y_{ij} = \mu_{ij} + Z_i + V_{ij},$$

where  $\mu_{ij} = EY_{ij}$  and  $Z_i \sim N(0, \nu^2)$  independent of  $V_{ij} \sim N(0, \tau^2)$  and it holds that  $\rho = \nu^2 / (\nu^2 + \tau^2)$  and  $\sigma^2 = \nu^2 + \tau^2$  (for  $\varepsilon_{ij} = Z_i + V_{ij}$ )

- **Interpretation:** linear model for the mean of the response with a random intercept (with the variance between subjects  $\nu^2 > 0$ )

# Exponential correlation model

- **Assumption:** covariance between  $Y_{ij}$  and  $Y_{ik}$  for  $i \neq k$  is of the form

$$v_{jk} = \sigma^2 \exp\{-\phi|t_j - t_k|\}$$

and it decays towards zero as the time separation between repeated observations increases (with the rate of decay given by  $\phi > 0$ )

- **Matrix notation:**  $\mathbb{V}_0 = (v_{jk})_{j,k=1}^n$
- **Motivation:** the response (random) variable  $Y_{ij}$  can be decomposed as

$$Y_{ij} = \mu_{ij} + W_{ij},$$

where  $W_{ij} = \rho W_{i(j-1)} + Z_{ij}$  for  $Z_{ij} \sim N(0, \sigma^2(1 - \rho^2))$  independent (verify, that it holds that  $\text{Var}Y_{ij} = \text{Var}W_{ij} = \sigma^2$ )

- **Interpretation:** linear model for the mean of the response with with the first order autoregressive correlation structure
- **Generalization:**  $Y_{ij} = \mu_{ij} + W_i(t_j)$  for continuous time Gaussian processes  $\{X_i(t); t \in \mathbb{R}\}$  independent for  $i = 1, \dots, N$  and general time points  $t_1 < \dots < t_{ni}$

## Towards least squares – two step estimation

- For simplification assume the model  $Y_{ij} = a + bX_{ij} + \varepsilon_{ij}$  and no distributional assumption for the error vector  $\varepsilon = (\varepsilon_{11}, \dots, \varepsilon_{Nn})^\top$
- Two-stage least squares as a simple estimation method for models where it is needed to distinguish the variance sources (within/between subjects)

## Towards least squares – two step estimation

- For simplification assume the model  $Y_{ij} = a + bX_{ij} + \varepsilon_{ij}$  and no distributional assumption for the error vector  $\varepsilon = (\varepsilon_{11}, \dots, \varepsilon_{Nn})^\top$
- Two-stage least squares as a simple estimation method for models where it is needed to distinguish the variance sources (within/between subjects)
- **Stage 1:** longitudinal profiles for each subject  $i \in \{1, \dots, N\}$  individually

$$Y_{ij} = A_i + B_i X_{ij} + W_{ij}, \quad j = 1, \dots, n, \quad \text{and } W_{ij} \sim (0, \tau^2), \quad i.i.d.$$

to obtain  $\hat{A}_i = A_i + Z_{ai}$  and  $\hat{B}_i = B_i + Z_{bi}$ , for  $Z_{ai} \sim (0, v_{ai}^2)$ ,  $Z_{bi} \sim (0, v_{bi}^2)$

- **Stage 2:** OLS analysis of the subject's specific parameter estimates

$$A_i = a + \delta_{ai} \quad \text{and} \quad B_i = b + \delta_{bi}$$

for independent errors  $\delta_{ai} \sim (0, \sigma_a^2)$  and  $\delta_{bi} \sim (0, \sigma_b^2)$

- **Therefore:**  $\hat{A}_i = a + (\delta_{ai} + Z_{ai})$  and  $\hat{B}_i = b + (\delta_{bi} + Z_{bi})$



# Summary

- Two alternative but not equivalent multivariate model formulations

$$\mathbf{Y} = \mathbf{X}\mathbf{B} + \mathbf{U} \quad \text{versus} \quad \mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

- Estimation of the unknown parameters in  $\mathbf{B} \in \mathbb{R}^{p \times n}$  or  $\boldsymbol{\beta} \in \mathbb{R}^p$  (either in terms of the least squares or the maximum likelihood estimation)
- Decomposition of the overall data variability into two different sources (the within subject's variability and the between subjects' variability)
- Marginal or hierarchical inference (in terms of the confidence intervals/regions or the statistical tests)
- Two stage estimation approach in the model  $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\beta}$  (towards the mixed effect model with fixed and random effects)