

1. Consider the data set `growthheight.txt` related to growth profiles of sample of boys and girls:

```
> head(data)
      gender  Y10   Y11   Y12   Y13   Y14   Y15   Y16   Y17   Y18
girl01  girl 138.6 146.8 153.1 156.2 157.7 158.2 158.6 158.7 158.9
girl02  girl 140.9 146.1 152.9 159.5 162.6 165.0 165.6 166.1 166.0
girl03  girl 148.8 155.6 159.6 160.3 161.6 161.7 161.9 161.7 162.2
girl04  girl 143.0 148.5 154.8 161.2 165.2 166.5 167.2 167.4 167.8
girl05  girl 141.0 147.0 153.0 161.0 166.0 168.0 169.0 170.0 170.0
> tail(data)
      gender  Y10   Y11   Y12   Y13   Y14   Y15   Y16   Y17   Y18
boy35   boy 154.0 160.9 166.3 171.0 177.3 183.4 186.2 186.7 188.0
boy36   boy 143.0 148.0 153.6 162.1 171.2 176.8 179.1 180.1 180.8
boy37   boy 148.3 154.2 161.8 171.1 177.9 181.3 182.2 183.1 183.7
boy38   boy 147.8 153.5 161.5 169.7 175.6 178.3 179.2 179.8 180.7
boy39   boy 139.8 145.1 150.2 156.7 164.1 171.0 174.7 176.1 176.4
```

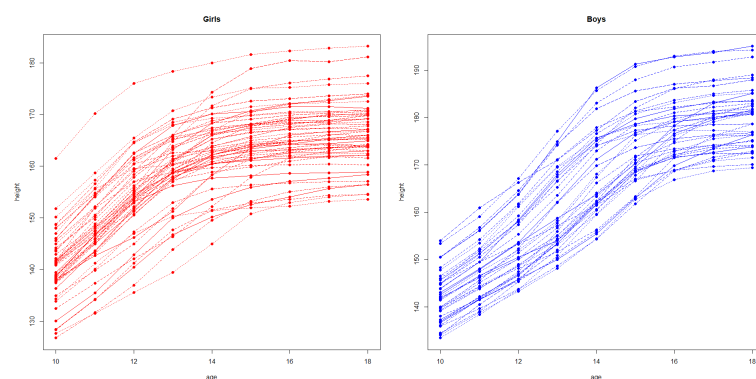
A list containing the heights of 39 boys and 54 girls from age 10 to 18 and the ages at which they were collected.

gender - values of boy and girl

Y10, Y11, ..., Y18 - the variables giving the heights in centimeters of children at ages 10,11,12,...18.

Tuddenham, R. D., and Snyder, M. M. (1954)

"Physical growth of California boys and girls from birth to age 18",  
University of California Publications in Child Development, 1, 183-364.



Denote variables as following  $Y_1 = Y_{10}, Y_2 = Y_{11}, \dots, Y_9 = Y_{18}$  and  $X =$  gender with index variable  $j$  associated to its values. Let us assume that the random vector (i.e, random vector for each row  $i$ )  $\mathbf{y}_i = (y_{i1}, y_{i2}, \dots, y_{i9})$  follows the normal distribution  $\mathbf{y}_i \sim N(\boldsymbol{\mu}_i, \boldsymbol{\Sigma})$ , where the expected value vector  $\boldsymbol{\mu}_i = (\mu_{i1}, \mu_{i2}, \dots, \mu_{i9})'$  is modeled by the growth curve model

$$\mathcal{M}: \quad \boldsymbol{\mu}_i = \begin{pmatrix} \mu_{i1} \\ \mu_{i2} \\ \mu_{i3} \\ \vdots \\ \mu_{i8} \\ \mu_{i9} \end{pmatrix} = \begin{pmatrix} 1 & t_1 & t_1^2 & t_1^3 \\ 1 & t_2 & t_2^2 & t_2^3 \\ 1 & t_3 & t_3^2 & t_3^3 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & t_8 & t_8^2 & t_8^3 \\ 1 & t_9 & t_9^2 & t_9^3 \end{pmatrix} \begin{pmatrix} \theta_{0j} \\ \theta_{1j} \\ \theta_{2j} \\ \theta_{3j} \end{pmatrix} = (\mathbf{1} : \mathbf{t} : \mathbf{t}^2 : \mathbf{t}^3) \begin{pmatrix} \theta_{0j} \\ \theta_{1j} \\ \theta_{2j} \\ \theta_{3j} \end{pmatrix} = \mathbf{T}\boldsymbol{\theta}_j,$$

where the time values are  $\mathbf{t} = (t_1, t_2, t_3, \dots, t_8, t_9)' = (10, 11, 12, \dots, 17, 18)'$ .

- (a) The model  $\mathcal{M}$  can be written for the whole data as  $\mathbf{Y} = \mathbf{X}\boldsymbol{\Theta}\mathbf{T}' + \mathbf{E}$ , where, for each row of  $\mathbf{E}$ , it is assumed  $\varepsilon_i \sim N(\mathbf{0}, \boldsymbol{\Sigma})$ , and  $\boldsymbol{\Theta} = \begin{pmatrix} \boldsymbol{\theta}'_1 \\ \boldsymbol{\theta}'_2 \end{pmatrix}$ . Find an estimate  $\hat{\boldsymbol{\Theta}}$  for the parameters  $\boldsymbol{\Theta}$ .

(1 point)

- (b) For the girls, find the estimate to the expected value  $\mu_{i10}$  at the age  $t_{10} = 19$ .

(1 point)

- (c) Predict the value of the new observation  $\mathbf{y}_f$  in both cases of a child being girl (i.e.,  $x_f = 1$ ) and a boy (i.e.,  $x_f = 0$ ). Create also a graph that displays the predicted profiles for a girl and a boy in the same graph.

(1 point)

- (d) Test at 5% significance level, is the explanatory variable  $X = \text{gender}$  statistically significant variable in the model. Calculate the value of the test statistic.

(1 point)

- (e) Test the hypotheses

$$H_0: \boldsymbol{\mu}_i = (\mathbf{1} : \mathbf{t}) \begin{pmatrix} \theta_{0j} \\ \theta_{1j} \end{pmatrix}, \quad H_1: \boldsymbol{\mu}_i = (\mathbf{1} : \mathbf{t} : \mathbf{t}^2 : \mathbf{t}^3) \begin{pmatrix} \theta_{0j} \\ \theta_{1j} \\ \theta_{2j} \\ \theta_{3j} \end{pmatrix}.$$

(1 point)

- (f) Consider the partitioned random vector

$$\mathbf{y}_f = \begin{pmatrix} \mathbf{y}_{f1} \\ \mathbf{y}_{f2} \end{pmatrix} = \begin{pmatrix} \mathbf{T}_1 \\ \mathbf{T}_2 \end{pmatrix} \boldsymbol{\Theta}' \mathbf{x}_f + \begin{pmatrix} \varepsilon_{f1} \\ \varepsilon_{f2} \end{pmatrix},$$

where  $\mathbf{y}_{f1}$  contains the random variables  $\mathbf{y}_{f1} = (y_{f1}, y_{f2}, y_{f3})'$ . Predict the value of the random vector  $\mathbf{y}_{f2} = (y_{f4}, y_{f5}, y_{f6}, y_{f7}, y_{f8}, y_{f9})'$  when

gender	Y10	Y11	Y12
girl	148.7	156.6	164.7

Create also 80 % simultaneous (asymptotic) prediction intervals for elements of the random vector  $\mathbf{y}_{f2}$ .

(1 point)

2. Consider the dataset `stageforest.txt`:

```
> data<-read.table("stageforest.txt", sep="\t", dec=".", header=TRUE)
> head(data)
  Tree.ID Age Forest.ID dbhib.cm height.m
1      1  55 Clearwater  37.084 21.76272
2      1  45 Clearwater  31.496 18.71472
3      1  35 Clearwater  22.352 12.22248
4      1  25 Clearwater  17.780  8.71728
5      1  15 Clearwater  10.160  5.97408
6      2 107 Clearwater  50.800 31.51632
> tail(data)
  Tree.ID Age Forest.ID dbhib.cm height.m
537     85  68  Wallowa  36.322 29.77896
538     85  58  Wallowa  31.750 28.98648
539     85  48  Wallowa  24.892 25.72512
540     85  38  Wallowa  17.526 18.31848
541     85  28  Wallowa  11.684 11.24712
542     85  18  Wallowa   6.604  6.18744
```

## Description

The data are stem measures from 66 trees.  
 The trees were selected as having been dominant throughout their lives with no visible evidence of damage or forks.  
 The trees came from stands throughout the inland range of the species.

A data frame with 542 observations on the following variables.

Tree.ID - A factor uniquely identifying the tree.

Age - Age of tree

Forest.ID - The national forest in which the tree was.

dbhib.cm - Diameter (cm.) at 1.37 m (4'6?).

height.m - Height of tree (m)

The national forests are: Kaniksu, Coeur d'Alene, St. Joe, Clearwater, Nez Perce, Clark Fork, Umatilla, Wallowa, and Payette.

Note that values are strangely in wrong order respect the age variable.

Denote the variables as following

$$Y_1 = \text{dbhib.cm}, \quad Y_2 = \text{height.m}, \quad T = \text{Age}.$$

For the tree  $i$ , consider the multivariate linear mixed effects model

$$\begin{aligned} \mathcal{M}: \quad y_{i1t} &= \beta_{01} + \beta_{11}t_i + \beta_{21}t_i^2 + b_{i01} + b_{i11}t + \varepsilon_{i1t}, \\ y_{i2t} &= \beta_{02} + \beta_{12}t_i + \beta_{22}t_i^2 + b_{i02} + b_{i12}t + \varepsilon_{i2t}, \end{aligned}$$

which can be written also as

$$\mathcal{M}: \quad \mathbf{Y}_i = \mathbf{X}_i \mathbf{B} + \mathbf{Z}_i \mathbf{R}_i + \mathbf{E}_i, \quad \text{vec}(\mathbf{E}_i) \sim N(\mathbf{0}, \Sigma \otimes \mathbf{I}),$$

where

$$\mathbf{Y}_i = (\mathbf{y}_{i1} : \mathbf{y}_{i2}), \quad \mathbf{X}_i = (\mathbf{1} : \mathbf{t}_i : \mathbf{t}_i^2), \quad \mathbf{Z}_i = (\mathbf{1} : \mathbf{t}_i), \quad \text{vec}(\mathbf{R}_i) \sim N(\mathbf{0}, \mathbf{F}).$$

(a) Calculate the estimate for the fixed parameters  $B$ . (2 points)

(b) Calculate the estimate for the covariance matrix  $\Sigma$ . (1 point)

(c) Predict the value of the new observation  $y_f = \begin{pmatrix} y_{f1t} \\ y_{f2t} \end{pmatrix}$ , when

Tree.ID=67, Age=97.

Create also 80 % simultaneous (asymptotic) prediction intervals for elements of the random vector  $y_f$ .

(2 points)

(d) Predict the value of the new observation  $y_{f2t}$ , i.e., predict the value of the variable  $Y_2 = \text{height.m}$  when we know that

Tree.ID=67, Age=97, dbhib.cm=35.5,

where the value dbhib.cm=35.5 means that at the age  $t = 97$  observed value  $y_{f1t} = 35.5$  is known.

(1 point)

3. (a) Consider the following longitudinal dataset:

```
> library(geepack)
> data(ohio)
> head(ohio)
  resp id age smoke
1     0  0 -2      0
2     0  0 -1      0
3     0  0  0      0
4     0  0  1      0
5     0  1 -2      0
6     0  1 -1      0
> tail(ohio)
  resp id age smoke
2143  1 535  0      1
2144  1 535  1      1
2145  1 536 -2      1
2146  1 536 -1      1
2147  1 536  0      1
2148  1 536  1      1
```

The ohio data frame has 2148 rows and 4 columns.

The dataset is a subset of the six-city study, a longitudinal study of the health effects of air pollution.

resp - an indicator of wheeze status (1=yes, 0=no)

id - a numeric vector for subject id

age - a numeric vector of age, 0 is 9 years old

smoke - an indicator of maternal smoking at the first year of the study

Denote the variables as following

$$Y = \text{resp}, \quad X = \text{smoke}, \quad T = \text{age},$$

with index  $i$  related to the values of  $\text{id}$ . Model the expected value of the random variable  $y_{it}$  by the interaction effect model

$$g(\mu_{it}) = \beta_0 + \beta_1 x_{it} + \beta_2 t + \beta_3 x_{it}t,$$

with taking into account that most likely, for each  $i$ , the values  $y_i$  are correlated. Calculate the estimate for the expected value  $\mu_{it}$  when

id	age	smoke
536	1.25	1

(3 points)

(b) In a growth curve model

$$\mathcal{M} : \mathbf{Y} = \mathbf{X}\boldsymbol{\Theta}\mathbf{T}' + \mathbf{E},$$

parameter matrix  $\boldsymbol{\Theta}$  can be estimated by the formula

$$\hat{\boldsymbol{\Theta}} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{Y}\mathbf{T}(\mathbf{T}'\mathbf{T})^{-1}.$$

The growth curve model  $\mathcal{M}$  can also be written in a form

$$\mathcal{V} : \text{vec}(\mathbf{Y}) = (\mathbf{T} \otimes \mathbf{X}) \text{vec}(\boldsymbol{\Theta}) + \text{vec}(\mathbf{E}).$$

Show that the ordinary least squares estimate for  $\text{vec}(\boldsymbol{\Theta})$

$$\min_{\text{vec}(\boldsymbol{\Theta})} [\text{vec}(\mathbf{Y}) - (\mathbf{T} \otimes \mathbf{X}) \text{vec}(\boldsymbol{\Theta})]' [\text{vec}(\mathbf{Y}) - (\mathbf{T} \otimes \mathbf{X}) \text{vec}(\boldsymbol{\Theta})]$$

is containing the same estimates as  $\hat{\boldsymbol{\Theta}}$ . Note that in a classical linear model  $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon}$ , the ordinary least squares estimate is  $\hat{\boldsymbol{\beta}} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}$ .

(3 points)