Homework 6

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Problem 1

Variance

$$Var(Y_{ij}) = Var(\mu + b_i + e_{ij})$$

$$= Var(\mu) + Var(b_i) + Var(e_{ij})$$

$$= \sigma_b^2 + \sigma_e^2$$

Covariance

$$\mu_{Y_{ij}} = \mu_{Y_{ik}} = E[\mu + b_i + e_{ij}] = \mu$$

As e_{ij} and e_{ik} are independent, $E[e_{ij}e_{ik}] = E[e_{ij}]E[e_{ik}] = 0$.

$$Cov(Y_{ij}, Y_{ik}) = E[(Y_{ij} - \mu_{ij})(Y_{ik} - \mu_{ik})]$$

$$= E[(b_i + e_{ij})(b_i + e_{ik})]$$

$$= E[b_i^2 + b_i(e_{ij} + e_{ik}) + e_{ij}e_{ik}]$$

$$= E[b_i^2] + E[b_i(e_{ij} + e_{ik})] + E[e_{ij}e_{ik}]$$

$$= Var(b_i) + [E(b_i)]^2 + b_i \times 0 + 0$$

$$= \sigma_b^2$$

Correlation

$$cor(Y_{ij}, Y_{ik}) = Cov(Y_{ij}, Y_{ik}) / \sqrt{var(Y_{ij}), var(Y_{ik})} = \frac{\sigma_b^2}{\sigma_b^2 + \sigma_e^2}$$

According to the correlation result, we can know that correlation between any two Y_{ij} are constant, so this correspond to compound symmetry covariance patterns.

Problem 2

```
library(ggplot2)
library(patchwork)
library(tidyverse)
library(nlme)
```

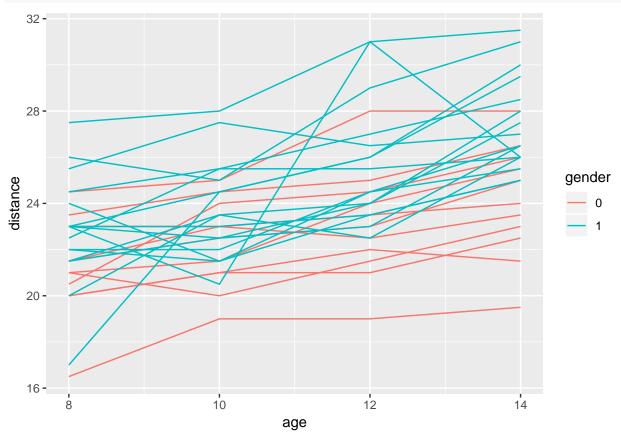
Question 1

Import dental data.

```
dental_data =
  read.csv("./HW6-dental.txt", sep="") %>%
  janitor::clean_names() %>%
  mutate(gender = as.factor(gender))
```

Make the spaghetti plot.

spag_p = ggplot(dental_data, aes(age, distance, group=child, color=gender)) + geom_line()
spag_p



Question 2

$$var(Yij) = var[\beta_0 + a_i + b_0 * I(sex_i = 0) + b_1 * I(sex_i = 1) + \beta_1 * age_{ij} + e_{ij}]$$

$$= var(a_i) + var(b_k) + var(e_{ij})$$

$$= \sigma_a^2 + \sigma_b^2 + \sigma_e^2$$

$$E(Y_{ij}) = \beta_0 + \beta_1 * age_{ij}$$

For the same individual in different ages:

$$cov(Y_{ij}, Y_{im}) = E[(a_i + b_k + e_{ij})(a_i + b_k + e_{im})]$$

$$= E[a_i^2 + a_i b_k + a_i e_{im} + a_i b_k + b_k^2 + b_k e_{im} + a_i e_{ij} + b_k e_{ij} + e_{ij} e_{im}]$$

$$= E(a_i^2) + E(b_k^2)$$

$$= \sigma_a^2 + \sigma_b^2$$

For different individuals:

1) same gender

When measured in different ages,

$$cov(Y_{hj}, Y_{im}) = E[(a_h + b_k + e_{hj})(a_i + b_k + e_{im})]$$

$$= E[a_i a_h + a_h b_k + a_h e_{im} + a_i b_k + b_k^2 + b_k e_{im} + a_i e_{hj} + b_k e_{hj} + e_{hj} e_{im}]$$

$$= E(b_k^2)$$

$$= \sigma_h^2$$

When measured in the same age,

$$cov(Y_{hj}, Y_{ij}) = E[(a_h + b_k + e_{hj})(a_i + b_k + e_{ij})]$$
$$= E(b_k^2)$$
$$= \sigma_h^2$$

2) different genders

$$cov(Y_{hj}, Y_{im}) = E[(a_h + b_0 + e_{hj})(a_i + b_1 + e_{im})]$$

= 0

So for same individuals:

$$M_s = \begin{bmatrix} \sigma_a^2 + \sigma_b^2 + \sigma_e^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 \\ \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 + \sigma_e^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 \\ \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 + \sigma_e^2 & \sigma_a^2 + \sigma_b^2 \\ \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 & \sigma_a^2 + \sigma_b^2 + \sigma_e^2 \end{bmatrix}$$

For different individuals:

$$M_d = \begin{bmatrix} \sigma_b^2 & \sigma_b^2 & \sigma_b^2 & \sigma_b^2 \\ \sigma_b^2 & \sigma_b^2 & \sigma_b^2 & \sigma_b^2 \\ \sigma_b^2 & \sigma_b^2 & \sigma_b^2 & \sigma_b^2 \\ \sigma_b^2 & \sigma_b^2 & \sigma_b^2 & \sigma_b^2 \end{bmatrix}$$

For all grils:

$$N_g = \begin{bmatrix} M_s & M_d & \dots & M_d \\ M_d & M_s & \dots & M_d \\ \vdots & \vdots & \ddots & \vdots \\ M_d & M_d & \dots & M_s \end{bmatrix}_{11 \times 11}$$

For all boys:

$$N_b = \begin{bmatrix} M_s & M_d & \dots & M_d \\ M_d & M_s & \dots & M_d \\ \vdots & \vdots & \ddots & \vdots \\ M_d & M_d & \dots & M_s \end{bmatrix}_{16 \times 16}$$

So the model in marginal form is:

$$E(Y_{ij}) = \beta_0 + a_i + b_0 * I(sex_i = 0) + b_1 * I(sex_i = 1) + \beta_1 * age_{ij} + e_{ij}$$

$$var(Y_i) = \begin{bmatrix} N_g & 0\\ 0 & N_b \end{bmatrix}_{27 \times 27}$$

Question 3

Compound symmetry covariance

```
comsym <- gls(distance~gender+age,dental_data,</pre>
              correlation=corCompSymm(form = ~ 1 | child), method="REML")
summary(comsym)
## Generalized least squares fit by REML
##
     Model: distance ~ gender + age
##
     Data: dental_data
##
          AIC
                    BIC
                           logLik
     447.5125 460.7823 -218.7563
##
##
## Correlation Structure: Compound symmetry
  Formula: ~1 | child
   Parameter estimate(s):
         Rho
##
## 0.6144914
##
## Coefficients:
                    Value Std.Error
                                     t-value p-value
## (Intercept) 15.385690 0.8959848 17.171820 0.0000
## gender1
                2.321023 0.7614169 3.048294 0.0029
                0.660185 0.0616059 10.716263 0.0000
## age
##
##
    Correlation:
           (Intr) gendr1
   gender1 -0.504
##
           -0.756 0.000
##
   age
##
## Standardized residuals:
##
                                    Med
                                                              Max
## -2.59712955 -0.64544226 -0.02540005 0.51680604 2.32947531
##
## Residual standard error: 2.305697
## Degrees of freedom: 108 total; 105 residual
According to the result given by R, the model with compound symmetry covariance is:
                           E(y_{ij}) = 15.396 + 2.151 * sex_i + 0.664 * age_{ij}
```

 $Var(Y_i) = is:$

```
corMatrix(comsym$modelStruct$corStruct)[[1]]*(comsym$sigma)^2
```

```
## [,1] [,2] [,3] [,4]
## [1,] 5.316240 3.266784 3.266784 3.266784
```

```
## [2,] 3.266784 5.316240 3.266784 3.266784
## [3,] 3.266784 3.266784 5.316240 3.266784
## [4,] 3.266784 3.266784 3.266784 5.316240
```

Exponential covariance

```
expo <- gls(distance~gender+age ,dental_data,</pre>
            correlation=corExp(form = ~ 1 | child), method="REML")
summary(expo)
## Generalized least squares fit by REML
##
     Model: distance ~ gender + age
##
     Data: dental data
          AIC
##
                    BIC
                           logLik
##
     455.4483 468.7181 -222.7241
##
## Correlation Structure: Exponential spatial correlation
## Formula: ~1 | child
   Parameter estimate(s):
##
      range
## 2.133938
##
## Coefficients:
##
                    Value Std.Error
                                      t-value p-value
## (Intercept) 15.459995 1.1309319 13.670138
                                                 0e+00
                                                 7e-04
  gender1
                2.418714 0.6933441
                                     3.488476
## age
                 0.652960 0.0906420 7.203723
                                                 0e+00
##
##
    Correlation:
           (Intr) gendr1
## gender1 -0.363
## age
           -0.882 0.000
##
## Standardized residuals:
##
           Min
                         Q1
                                    Med
                                                  QЗ
                                                              Max
## -2.65148774 -0.69592567 -0.06214639 0.48659340 2.29666951
##
## Residual standard error: 2.301495
## Degrees of freedom: 108 total; 105 residual
According to the result given by R, the model with exponential covariance is:
                           E(y_{ij}) = 15.460 + 2.419 * sex_i + 0.653 * age_{ij}
Var(Y_i) = is:
corMatrix(expo$modelStruct$corStruct)[[1]]*(expo$sigma)^2
            [,1]
                      [,2]
                               [,3]
## [1,] 5.296881 3.315144 2.074839 1.298574
## [2,] 3.315144 5.296881 3.315144 2.074839
## [3,] 2.074839 3.315144 5.296881 3.315144
## [4,] 1.298574 2.074839 3.315144 5.296881
```

Autoregressive covariance

```
auto1 <- gls(distance~gender+age ,dental_data,</pre>
              correlation=corAR1(form = ~ 1 | child), method="REML")
summary(auto1)
## Generalized least squares fit by REML
##
     Model: distance ~ gender + age
##
     Data: dental_data
##
          AIC
                    BIC
                           logLik
##
     455.4483 468.7181 -222.7241
##
## Correlation Structure: AR(1)
    Formula: ~1 | child
##
##
    Parameter estimate(s):
##
         Phi
##
  0.6258671
##
## Coefficients:
##
                                       t-value p-value
                    Value Std.Error
## (Intercept) 15.459995 1.1309319 13.670138
                                                  0e+00
                                                  7e-04
   gender1
                 2.418714 0.6933441 3.488476
## age
                 0.652960 0.0906420 7.203723
                                                  0e+00
##
##
    Correlation:
##
            (Intr) gendr1
  gender1 -0.363
##
           -0.882 0.000
  age
##
## Standardized residuals:
##
           Min
                         Q1
                                                   QЗ
                                                               Max
                                     Med
  -2.65148770 -0.69592566 -0.06214639
                                          0.48659339
##
## Residual standard error: 2.301495
## Degrees of freedom: 108 total; 105 residual
According to the result given by R, the model with autoregressive covariance is:
                           E(y_{ij}) = 15.460 + 2.419 * sex_i + 0.653 * age_{ij}
Var(Y_i) = is:
corMatrix(auto1$modelStruct$corStruct)[[1]]*(auto1$sigma)^2
##
             [,1]
                      [,2]
                                [,3]
                                          [,4]
## [1,] 5.296881 3.315144 2.074840 1.298574
## [2,] 3.315144 5.296881 3.315144 2.074840
```

Conclusion:

[3,] 2.074840 3.315144 5.296881 3.315144 ## [4,] 1.298574 2.074840 3.315144 5.296881

According to results given by R, models with exponential covariance and autoregressive covariance have same coefficient parameter estimates and covariance estimates as model with exponential covariance is a generalization of autoregressive pattern and in dental data, measurements made at equal intervals of time.

The model with compound symmetry covariance have different coefficient parameter estimates and covariance estimates with other two models but the parameter estimates are closed. Since models with compound symmetry assume correlation between any two visits are constant while other two models do not have such assumption, their covariance estimates are different.