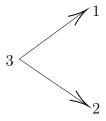
GENE638 - Homework 5

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Data:

| COW | YEAR | TICK COUNT |
|-----|------|------------|
| 1 | 1 | 80 |
| 2 | 1 | 70 |
| 1 | 2 | 86 |
| 2 | 2 | 72 |
| 1 | 3 | 92 |
| 2 | 3 | 78 |

Pedrigree:



1. Construct the numerator relationship and its inverse

$$A = \begin{bmatrix} 1.00 & 0.25 & 0.50 \\ 0.25 & 1.00 & 0.50 \\ 0.50 & 0.50 & 1.00 \end{bmatrix} \qquad A^{-1} = \begin{bmatrix} 1.33 & 0.00 & -0.67 \\ 0.00 & 1.33 & -0.67 \\ -0.67 & -0.67 & 1.67 \end{bmatrix}$$

2. The predicted breeding value of the unidentified parent of animal 1 is $\frac{2}{3}(\hat{A}_1 - \frac{1}{2}\hat{A}_3)$ and that for the unidentified parent of animal 2 is $\frac{2}{3}(\hat{A}_2 - \frac{1}{2}\hat{A}_3)$. Identify the base generation animals and show that $1'A^{-1}\hat{u} = 0$ is consistent with the sum of breeding values of base generation being 0.

$$\begin{bmatrix} 1 & 1 & 1 & 1 & 1 \end{bmatrix} \begin{bmatrix} 2.00 & 0.00 & -1.00 & -1.00 & -0.00 \\ 0.00 & 2.00 & -1.00 & -0.00 & -1.00 \\ -1.00 & -1.00 & 2.00 & 0.50 & 0.50 \\ -1.00 & -0.00 & 0.50 & 1.50 & 0.00 \\ -0.00 & -1.00 & 0.50 & 0.00 & 1.50 \end{bmatrix} \begin{bmatrix} 4.38 \\ -4.38 \\ 0.00 \\ 2.92 \\ -2.92 \end{bmatrix} = -1.332268e - 15$$

$$\begin{bmatrix} A1 & A2 & A3 & M1 & M2 \\ 5.84 & -5.84 & 0.00 & 0.00 & 0.00 \end{bmatrix}$$

3. Write the data in general matrix terms $y = X\beta + Zu + e$ for the model $Y_{ij} = \text{Year}_i + \text{Animal}_j + e_{ij}$ where $\text{Year}_i = \mu + \text{Year}_i$. Include animal 3 in \hat{u} as in (2)

$$\begin{bmatrix} 80 \\ 70 \\ 86 \\ 72 \\ 92 \\ 78 \end{bmatrix} = \begin{bmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{bmatrix} + \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \end{bmatrix} \begin{bmatrix} u_1 \\ u_2 \\ u_3 \end{bmatrix} + \begin{bmatrix} e_{11} \\ e_{21} \\ e_{12} \\ e_{22} \\ e_{13} \\ e_{23} \end{bmatrix}$$

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4. Assuming $R = I\sigma_e^2$ and $G = A\sigma_a^2$ and $\lambda = \frac{\sigma_e^2}{\sigma_a^2} = 3$, write the MME for model in (3)

$$\begin{bmatrix} 2 & 0 & 0 & 1 & 1 & 0 \\ 0 & 2 & 0 & 1 & 1 & 0 \\ 0 & 0 & 2 & 1 & 1 & 0 \\ 1 & 1 & 1 & 7 & 0 & -2 \\ 1 & 1 & 1 & 0 & 7 & -2 \\ 0 & 0 & 0 & -2 & -2 & 5 \end{bmatrix} \begin{bmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \\ u_1 \\ u_2 \\ u_3 \end{bmatrix} = \begin{bmatrix} 150 \\ 158 \\ 170 \\ 258 \\ 220 \\ 0 \end{bmatrix}$$

5. From the row of the MME corresponding to \hat{A}_3 show that $\hat{A}_3 = \frac{2}{5}(\hat{A}_1 + \hat{A}_2)$

$$0 = -2\hat{A}_1 - 2\hat{A}_2 + 5\hat{A}_3$$
$$-5\hat{A}_3 = -2\hat{A}_1 - 2\hat{A}_2$$
$$5\hat{A}_3 = 2\hat{A}_1 + 2\hat{A}_2$$
$$\hat{A}_3 = \frac{2\hat{A}_1 + 2\hat{A}_2}{5}$$
$$\hat{A}_3 = \frac{2}{5}(\hat{A}_1 + \hat{A}_2)$$

6. Absorb the year equations into the animal equations to obtain a system of equations involving only the unknown breeding values \hat{A}_1 , \hat{A}_2 , and \hat{A}_3

$$\begin{bmatrix} 0.280 & 0.137 & 0.167 \\ 0.137 & 0.280 & 0.167 \\ 0.167 & 0.167 & 0.333 \end{bmatrix} \begin{bmatrix} u_1 \\ u_2 \\ u_3 \end{bmatrix} = \begin{bmatrix} 19.0 \\ -19.0 \\ 0.0 \end{bmatrix}$$

7. What are the effective numbers of observations on all three animals?

8. From the appropriate row of the absorbed MME, once again show that $\hat{A}_3 = \frac{2}{5}(\hat{A}_1 + \hat{A}_2)$

$$C_{22} = \begin{bmatrix} 5.5 & -1.5 & -2.0 \\ -1.5 & 5.5 & -2.0 \\ -2.0 & -2.0 & 5.0 \end{bmatrix}$$

$$-5\hat{A}_3 = -2\hat{A}_1 - 2\hat{A}_2$$

$$5\hat{A}_3 = 2\hat{A}_1 + 2\hat{A}_2$$

$$\hat{A}_3 = \frac{2\hat{A}_1 + 2\hat{A}_2}{5}$$

$$\hat{A}_3 = \frac{2}{5}(\hat{A}_1 + \hat{A}_2)$$

 $0 = -2\hat{A}_1 - 2\hat{A}_2 + 5\hat{A}_3$

9. Using ordinary Gauss-Seidel iteration, find two succesive approximations to the predicted breeding values in (6)

$$L = \begin{bmatrix} 0.0 & 0.0 & 0.0 \\ -1.5 & 0.0 & 0.0 \\ -2.0 & -2.0 & 0.0 \end{bmatrix} \qquad D = \begin{bmatrix} 5.5 & 0.0 & 0.0 \\ 0.0 & 5.5 & 0.0 \\ 0.0 & 0.0 & 5.0 \end{bmatrix} \qquad b = \begin{bmatrix} 19.0 \\ -19.0 \\ 0.0 \end{bmatrix}$$

```
> L \leftarrow D \leftarrow matrix(0, nrow = nrow(C22), ncol = ncol(C22))
> L[lower.tri(L)] <- C22[lower.tri(C22)]</pre>
> diag(D) <- diag(C22)
> x0 <- solve(D) %*% b
> for(i in 1:2){
    Xi \leftarrow solve(L + D) %*% (b - t(L) %*% x0)
    rownames(Xi) \leftarrow c("A1", "A2", "A3")
    print(Xi)
    x0 <- Xi
+ }
          [,1]
A1 2.5123967
A2 -2.7693464
A3 -0.1027799
    2.6618947
A2 -2.7659487
A3 -0.0416216
```

10. Show that $\hat{A}_1 = 2.7143$, $\hat{A}_2 = -2.7143$ and $\hat{A}_3 = 0$, provide a solution to the absorbed equations in (6)

$$\begin{bmatrix} 0.280 & 0.137 & 0.167 \\ 0.137 & 0.280 & 0.167 \\ 0.167 & 0.167 & 0.333 \end{bmatrix} \begin{bmatrix} 2.7143 \\ -2.7143 \\ 0 \end{bmatrix} = \begin{bmatrix} 19.0 \\ -19.0 \\ 0.0 \end{bmatrix}$$

11. Using the breeding values in (10), backsolve the solutions to \hat{Y}_1 , \hat{Y}_2 and \hat{Y}_3

> beta <- solve(
$$t(X) \%*\% X) \%*\% t(X)\%*\%(y - Z \%*\% u)$$
 > beta

12. The inverse of the coefficient matrix in (6) for animals in the order \hat{A}_3 , \hat{A}_1 , \hat{A}_2 , is

$$\begin{bmatrix} 0.33330 & 0.16667 & 0.16667 \\ 0.16667 & 0.27976 & 0.13690 \\ 0.16667 & 0.1369 & 0.27976 \end{bmatrix} = (Z'MZ + A^{-1}\lambda)^{-1}$$

Which also is a submatrix of the inverse

$$\left[\begin{array}{cc} X'X & X'Z \\ Z'X & Z'Z + A^{-1}\lambda \end{array}\right] \left[\begin{array}{cc} C_{11} & C_{12} \\ C_{21} & C_{22} \end{array}\right]$$

Calculate the first approximations:

$$\sigma_e^2 = \frac{y'y - \hat{\underline{\beta}}'X'y - \hat{u}'Z'\underline{y}}{N - p}$$

$$\sigma_a^2 = \frac{\hat{u}'A^{-1}\hat{u} + \sigma_e^2 tr\left[A^{-1}C_{22}\right]}{q}$$

$$\lambda = \frac{\sigma_e^2}{\sigma_a^2}$$

```
> N <- length(y)</pre>
> p <- Matrix::rankMatrix(X)</pre>
> q \leftarrow ncol(Z)
> sigma2E <-(t(y) %*% y - t(beta) %*% t(X) %*%
                y - t(u) %*% t(Z) %*% y) / as.numeric(N - p)
> sigma2E
         [,1]
[1,] 47.61905
> sigma2A <- (t(u) %*% Ainv %*% u +
                 sigma2E * sum(diag(Ainv %*% solve(C22))))/q
> sigma2A
        [,1]
[1,] 20.1542
> sigma2E/sigma2A
         [,1]
[1,] 2.362736
```