

# Livestock Breeding and Genomics - Solution 11

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*2019-12-06*

## Problem 1 Multivariate BLUP Animal Model

The table below contains data for pre-weaning gain (WWG) and post-weaning gain (PWG) for 5 beef calves.

Animal	Sex	Sire	Dam	WWG	PWG
4	Male	1	NA	4.5	6.8
5	Female	3	2	2.9	5.0
6	Female	1	2	3.9	6.8
7	Male	4	5	3.5	6.0
8	Male	3	6	5.0	7.5

The genetic variance-covariance matrix  $G_0$  between the traits is

$$G_0 = \begin{bmatrix} 20 & 18 \\ 18 & 40 \end{bmatrix}$$

The residual variance-covariance matrix  $R_0$  between the traits is

$$R_0 = \begin{bmatrix} 40 & 11 \\ 11 & 30 \end{bmatrix}$$

### Your Task

Set up the mixed model equations for a multivariate BLUP analysis and compute the estimates for the fixed effects and the predictions for the breeding values.

### Solution

The matrices  $X_1$  and  $X_2$  relate records of PWG and WWG to sex effects. For both traits, we have an effect for the male and female sex. Hence the vector  $\beta$  of fixed effects corresponds to

$$\beta = \begin{bmatrix} \beta_{M,WWG} \\ \beta_{F,WWG} \\ \beta_{M,PWG} \\ \beta_{F,PWG} \end{bmatrix}$$

The matrices  $X_1$  and  $X_2$  are the same and correspond to

$$X_1 = X_2 = \begin{bmatrix} 1 & 0 \\ 0 & 1 \\ 0 & 1 \\ 1 & 0 \\ 1 & 0 \end{bmatrix}$$

Combining them to the multivariate version leads to

$$X = \begin{bmatrix} X_1 & 0 \\ 0 & X_2 \end{bmatrix}$$

$$X = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

Using the matrix  $X$  together with matrix  $R = I_n \otimes R_0$  to get

$$X^T R^{-1} X = \begin{bmatrix} 0.083 & 0.000 & -0.031 & 0.000 \\ 0.000 & 0.056 & 0.000 & -0.020 \\ -0.031 & 0.000 & 0.111 & 0.000 \\ 0.000 & -0.020 & 0.000 & 0.074 \end{bmatrix}$$

Similarly to the fixed effects, we can put together the vector of breeding values  $a$ .

$$a = \begin{bmatrix} a_{1,WWG} \\ a_{2,WWG} \\ a_{3,WWG} \\ a_{4,WWG} \\ a_{5,WWG} \\ a_{6,WWG} \\ a_{7,WWG} \\ a_{8,WWG} \\ a_{1,PWG} \\ a_{2,PWG} \\ a_{3,PWG} \\ a_{4,PWG} \\ a_{5,PWG} \\ a_{6,PWG} \\ a_{7,PWG} \\ a_{8,PWG} \end{bmatrix}$$

The design matrices  $Z_1$  and  $Z_2$  are equal and they link observations to breeding values.

$$Z_1 = Z_2 = \begin{bmatrix} 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{bmatrix}$$

$$Z = \begin{bmatrix} Z_1 & 0 \\ 0 & Z_2 \end{bmatrix}$$

$$Z = \begin{bmatrix} 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{bmatrix}$$

Together with the numerator relationship matrix  $A$  we can get  $G = G_0 \otimes A$  and from this  $G^{-1} = G_0^{-1} \otimes A^{-1}$

$$A^{-1} = \begin{bmatrix} 1.833 & 0.500 & 0.000 & -0.667 & 0.000 & -1.000 & 0.000 & 0.000 \\ 0.500 & 2.000 & 0.500 & 0.000 & -1.000 & -1.000 & 0.000 & 0.000 \\ 0.000 & 0.500 & 2.000 & 0.000 & -1.000 & 0.500 & 0.000 & -1.000 \\ -0.667 & 0.000 & 0.000 & 1.833 & 0.500 & 0.000 & -1.000 & 0.000 \\ 0.000 & -1.000 & -1.000 & 0.500 & 2.500 & 0.000 & -1.000 & 0.000 \\ -1.000 & -1.000 & 0.500 & 0.000 & 0.000 & 2.500 & 0.000 & -1.000 \\ 0.000 & 0.000 & 0.000 & -1.000 & -1.000 & 0.000 & 2.000 & 0.000 \\ 0.000 & 0.000 & -1.000 & 0.000 & 0.000 & -1.000 & 0.000 & 2.000 \end{bmatrix}$$

$$G^{-1} = \begin{bmatrix} 0.15 & 0.04 & 0.00 & -0.06 & 0.00 & -0.08 & 0.00 & 0.00 & -0.07 & -0.02 & 0.00 & 0.03 & 0.00 & 0.04 & 0.00 & 0.00 \\ 0.04 & 0.17 & 0.04 & 0.00 & -0.08 & -0.08 & 0.00 & 0.00 & -0.02 & -0.08 & -0.02 & 0.00 & 0.04 & 0.04 & 0.00 & 0.00 \\ 0.00 & 0.04 & 0.17 & 0.00 & -0.08 & 0.04 & 0.00 & -0.08 & 0.00 & -0.02 & -0.08 & 0.00 & 0.04 & -0.02 & 0.00 & 0.04 \\ -0.06 & 0.00 & 0.00 & 0.15 & 0.04 & 0.00 & -0.08 & 0.00 & 0.03 & 0.00 & 0.00 & -0.07 & -0.02 & 0.00 & 0.04 & 0.00 \\ 0.00 & -0.08 & -0.08 & 0.04 & 0.21 & 0.00 & -0.08 & 0.00 & 0.00 & 0.04 & 0.04 & -0.02 & -0.09 & 0.00 & 0.04 & 0.00 \\ -0.08 & -0.08 & 0.04 & 0.00 & 0.00 & 0.21 & 0.00 & -0.08 & 0.04 & 0.04 & -0.02 & 0.00 & 0.00 & -0.09 & 0.00 & 0.04 \\ 0.00 & 0.00 & 0.00 & -0.08 & -0.08 & 0.00 & 0.17 & 0.00 & 0.00 & 0.00 & 0.00 & 0.04 & 0.04 & 0.00 & -0.08 & 0.00 \\ 0.00 & 0.00 & -0.08 & 0.00 & 0.00 & -0.08 & 0.00 & 0.17 & 0.00 & 0.00 & 0.04 & 0.00 & 0.00 & 0.04 & 0.00 & -0.08 \\ -0.07 & -0.02 & 0.00 & 0.03 & 0.00 & 0.04 & 0.00 & 0.00 & 0.08 & 0.02 & 0.00 & -0.03 & 0.00 & -0.04 & 0.00 & 0.00 \\ -0.02 & -0.08 & -0.02 & 0.00 & 0.04 & 0.04 & 0.00 & 0.00 & 0.02 & 0.08 & 0.02 & 0.00 & -0.04 & -0.04 & 0.00 & 0.00 \\ 0.00 & -0.02 & -0.08 & 0.00 & 0.04 & -0.02 & 0.00 & 0.04 & 0.00 & 0.02 & 0.08 & 0.00 & -0.04 & 0.02 & 0.00 & -0.04 \\ 0.03 & 0.00 & 0.00 & -0.07 & -0.02 & 0.00 & 0.04 & 0.00 & -0.03 & 0.00 & 0.00 & 0.08 & 0.02 & 0.00 & -0.04 & 0.00 \\ 0.00 & 0.04 & 0.04 & -0.02 & -0.09 & 0.00 & 0.04 & 0.00 & 0.00 & -0.04 & -0.04 & 0.02 & 0.11 & 0.00 & -0.04 & 0.00 \\ 0.04 & 0.04 & -0.02 & 0.00 & 0.00 & -0.09 & 0.00 & 0.04 & -0.04 & -0.04 & 0.02 & 0.00 & 0.00 & 0.11 & 0.00 & -0.04 \\ 0.00 & 0.00 & 0.00 & 0.04 & 0.04 & 0.00 & -0.08 & 0.00 & 0.00 & 0.00 & 0.00 & -0.04 & -0.04 & 0.00 & 0.08 & 0.00 \\ 0.00 & 0.00 & 0.04 & 0.00 & 0.00 & 0.04 & 0.00 & -0.08 & 0.00 & 0.00 & -0.04 & 0.00 & 0.00 & -0.04 & 0.00 & 0.08 \end{bmatrix}$$

Using the matrices  $X$ ,  $Z$ ,  $R^{-1}$  and  $G^{-1}$ , we can compute  $Z^T R^{-1} X$  and  $Z^T R^{-1} Z + G^{-1}$ . These matrices define the right-hand side of the mixed model equations. But they are too big to be shown here.

The vector  $y$  of observations contains all observations of both traits

$$y = \begin{bmatrix} 4.50 \\ 2.90 \\ 3.90 \\ 3.50 \\ 5.00 \\ 6.80 \\ 5.00 \\ 6.80 \\ 6.00 \\ 7.50 \end{bmatrix}$$

The right-hand side is computed as

$$\begin{bmatrix} X^T R^{-1} y \\ Z^T R^{-1} y \end{bmatrix}$$

The solutions are

$$\begin{bmatrix} \widehat{\beta_{M,WWG}} \\ \widehat{\beta_{F,WWG}} \\ \widehat{\beta_{M,PWG}} \\ \widehat{\beta_{F,PWG}} \\ \widehat{a_{1,WWG}} \\ \widehat{a_{2,WWG}} \\ \widehat{a_{3,WWG}} \\ \widehat{a_{4,WWG}} \\ \widehat{a_{5,WWG}} \\ \widehat{a_{6,WWG}} \\ \widehat{a_{7,WWG}} \\ \widehat{a_{8,WWG}} \\ \widehat{a_{1,PWG}} \\ \widehat{a_{2,PWG}} \\ \widehat{a_{3,PWG}} \\ \widehat{a_{4,PWG}} \\ \widehat{a_{5,PWG}} \\ \widehat{a_{6,PWG}} \\ \widehat{a_{7,PWG}} \\ \widehat{a_{8,PWG}} \end{bmatrix} = \begin{bmatrix} 4.3609 \\ 3.3973 \\ 6.7999 \\ 5.8803 \\ 0.1509 \\ -0.0154 \\ -0.0784 \\ -0.0102 \\ -0.2703 \\ 0.2758 \\ -0.3161 \\ 0.2438 \\ 0.2796 \\ -0.0076 \\ -0.1703 \\ -0.0127 \\ -0.4778 \\ 0.5172 \\ -0.4790 \\ 0.3920 \end{bmatrix}$$

## Problem 2 Comparison of Reliabilites

Compare the predicted breeding values and the reliabilites obtained as results of Problem 1 with results from two univariate analyses for the same traits are used in Problem 1. All parameters can be taken from Problem 1.

### Solution