Additional Aspects of BLUP

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Aspects

- Accurracy
  - Results from MME are estimates of fixed effects and predictions of breeding values
  - Need statement about quality of estimates and predictions
- Confidence Intervals
- Decomposition of Predicted Breeding values

Accurracy

- One property of BLUP was that variance of prediction error is minimal
- How can we measure the variance of the prediction error
- Fixed effects

$$var(\beta - \hat{\beta}) = var(\hat{\beta})$$

- Random effects

$$var(u - \hat{u}) = var(u) - 2 * cov(u, \hat{u}) + var(\hat{u}) = var(u) - var(\hat{u}) = PEV(\hat{u})$$

because with BLUP: $cov(u, \hat{u}) = var(\hat{u})$

PEV

- PEV depends on inverse of coefficient matrix of MME

$$\left[\begin{array}{lr} X^T R^{-1} X & X^T R^{-1} Z \\ Z^T R^{-1} X & Z^T R^{-1} Z + G^{-1} \end{array}\right]^{-1} = \left[\begin{array}{cc} C^{11} & C^{12} \\ C^{21} & C^{22} \end{array}\right]$$

- For predicted breeding values $\hat{u}$

$$PEV(\hat{u}) = var(u) - var(\hat{u}) = C^{22}$$

Single Animal i

$$PEV(\hat{u}_i) = (C)_{ii}^{22}$$

where $(C)_{ii}^{22}$ is the i-th diagonal of C^{22}

- Accuracy measured by correlation

$$r_{u_i,\hat{u}_i} = \frac{cov(u_i, \hat{u}_i)}{\sqrt{var(u_i) * var(\hat{u}_i)}} = \sqrt{\frac{var(\hat{u}_i)}{var(u_i)}}$$

- Combining

$$PEV(\hat{u}_i) = (C)_{ii}^{22} = var(u_i) - var(\hat{u}_i) = var(u_i) - r_{u_i,\hat{u}_i}^2 var(u_i)$$

Accuracy B_i

$$B_i = r^2_{u_i,\hat{u}_i} = \frac{var(u_i) - (C)^{22}_{ii}}{var(u_i)} = 1 - \frac{PEV(\hat{u}_i)}{var(u_i)} = 1 - \frac{(C)^{22}_{ii}}{var(u_i)}$$

- B_i is large for small $PEV(\hat{u}_i)$
- In the limit $B_i \rightarrow 1$ for $PEV(\hat{u}_i) \rightarrow 0$
- For $PEV(\hat{u}_i) \rightarrow 0$ we must have $var(\hat{u}_i) \rightarrow var(u_i)$
- Therefore, the closer $var(\hat{u}_i)$ is to $var(u_i)$, the more accurate the predicted breeding value

Confidence Intervals of $\hat{u}_i$

- Predicted breeding value ($\hat{u}_i$) is a function of the data (y)
- Hence $\hat{u}_i$ is a random variable with a distribution

Standard Error of Prediction (SEP)

Widths Of Confidence Intervals

Table 1: Widths of Confidence Intervals for Given Accuracies

Accurracy	Interval Width
0.40	36.44
0.50	33.26
0.60	29.75
0.70	25.76
0.80	21.04
0.90	14.88
0.95	10.52
0.99	4.70

with $\hat{u}_i = 100$, $var(u_i) = 144$ and $\alpha = 0.05$

Decomposition of Predicted Breeding Value

- Write MME as

$$M \cdot s = r$$

with

$$s = \left[\begin{array}{c} \hat{\beta} \\ \hat{u} \end{array}\right]$$

- $\hat{\beta}$ has length p
- $\hat{u}$ has length q

Simplified Model

$$y_i = \mu + u_i + e_i$$

where

y_i	Observation for animal i
u_i	breeding value of animal i with a variance of $(1 + F_i)\sigma_u^2$
e_i	random residual effect with variance σ_e^2
μ	single fixed effect

Data

- all animals have an observation
- animal i has
  - parents s and d
  - n progeny k_j (with $j = 1, \ldots, n$)
  - n mates l_j (with $j = 1, \ldots, n$).
- progeny k_j has parents i and l_j.

Example

Animal	Sire	Dam	WWG
1	NA	NA	4.5
2	NA	NA	2.9
3	NA	NA	3.9
4	1	2	3.5
5	4	3	5.0

Variance components $\sigma_e^2 = 40$ and $\sigma_u^2 = 20$.

Model Components

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

$$X^T X = \begin{bmatrix} 5 \end{bmatrix},\ X^T Z = \begin{bmatrix} 1 & 1 & 1 & 1 & 1 \end{bmatrix}$$

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

Right-hand Side

$$X^T y = \left[\sum_{j=1}^{n} y_i \right] = 19.8$$

$$Z^T y = \begin{bmatrix} y1 \\ y2 \\ y3 \\ y4 \\ y5 \end{bmatrix} = \begin{bmatrix} 4.5 \\ 2.9 \\ 3.9 \\ 3.5 \\ 5 \end{bmatrix}$$

A^{-1}

$$A^{-1} = \begin{bmatrix} 1.5 & 0.5 & 0 & -1 & 0 \\ 0.5 & 1.5 & 0 & -1 & 0 \\ 0 & 0 & 1.5 & 0.5 & -1 \\ -1 & -1 & 0.5 & 2.5 & -1 \\ 0 & 0 & -1 & -1 & 2 \end{bmatrix}$$

MME

$$\begin{bmatrix} X^T X & X^T Z \\ Z^T X & Z^T Z + A^{-1} * \lambda \end{bmatrix} \begin{bmatrix} \hat{\mu} \\ \hat{u} \end{bmatrix} = \begin{bmatrix} X^T y \\ Z^T y \end{bmatrix}$$

Insert Data

$$\begin{bmatrix} 5 & 1 & 1 & 1 & 1 & 1 \\ 1 & 4 & 1 & 0 & -2 & 0 \\ 1 & 1 & 4 & 0 & -2 & 0 \\ 1 & 0 & 0 & 4 & 1 & -2 \\ 1 & -2 & -2 & 1 & 6 & -2 \\ 1 & 0 & 0 & -2 & -2 & 5 \end{bmatrix} \begin{bmatrix} \mu \\ \hat{u}1 \\ \hat{u}2 \\ \hat{u}3 \\ \hat{u}4 \\ \hat{u}5 \end{bmatrix} = \begin{bmatrix} 19.8 \\ 4.5 \\ 2.9 \\ 3.9 \\ 3.5 \\ 5 \end{bmatrix}$$

Animal 4

- parents 1 and 2
- progeny 5
- mate 3
- inspection of second but last equation in MME where y_4 and $\hat{u}_4$ occur
- Remember from construction of A^{-1}, the variable d^{ii} can assume the following values

$$d^{ii} = \begin{cases} 2 & \text{both parents known} \\ \frac{4}{3} & \text{one parent known} \\ 1 & \text{both parents unknown} \end{cases}$$

Extract Equation

$$y_4 = 3.5 = 1 * \hat{\mu} - 2 * \hat{u}_1 - 2 * \hat{u}_2 + 1 * \hat{u}_3 + 6 * \hat{u}_4 - 2 * \hat{u_5}$$

- Solving for $\hat{u}_4$

$$\hat{u}_4 = \frac{1}{6} \left[y_4 - \hat{\mu} + 2 * (\hat{u}_1 + \hat{u}_2) - \hat{u}_3 + 2\hat{u}_5 \right]$$

- $\hat{u}_4$ depends on
  - own performance record y_4
  - estimate of fixed effect $\hat{\mu}$ - environment
  - predicted breeding value of parents 1 and 2, mate 3 and progeny 5

General Equation

$$\hat{u}_i = \frac{1}{1 + \alpha\delta^{(i)} + \frac{\alpha}{4}\sum_{j=1}^{n}\delta^{(k_j)}} \Bigg[y_i - \hat{\mu} + \frac{\alpha}{2}\left\{ \delta^{(i)}(\hat{u}_s + \hat{u}_d) + \sum_{j=1}^{n}\delta^{(k_j)}(\hat{u}_{k_j} - \frac{1}{2}\hat{u}_{l_j}) \right\} \Bigg]$$

where	α	ration between variance components σ_e^2/σ_u^2
	$\delta^{(j)}$	contribution for animal j to A^{-1}