

OHP Picture 1

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Recap 2023-12-01

- Prediction of breeding values using BLUP-animal model
- BLUP animal model: Linear mixed effects model (LME)

$$y = X\beta + Zu + e$$

with β : fixed effects
 u : breeding values
 y : response variable or observations
 e : residuals

- BLUE-estimates $\hat{\beta}$ for fixed effects β
 BLUP-predictions \hat{u} for breeding values u
- Mixed model equations:

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

$\lambda = \frac{\sigma_e^2}{\sigma_u^2}$
 A^{-1} : inverse numerator relationship matrix

$$M \cdot \hat{s} = r$$

$$\hat{s} = \begin{bmatrix} \hat{\beta} \\ \hat{u} \end{bmatrix} = M^{-1} \cdot r \Rightarrow \text{solve}(M, r) \text{ in } R$$

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BLUE of β and BLUP of u have the following properties:

- Linear function of y
- Unbiased $\Rightarrow E[\hat{\beta}] = E[\beta]$; $E[\hat{u}] = E[u]$
- Best: $\text{var}(\beta - \hat{\beta})$ and $\text{var}(u - \hat{u})$ minimal
 - $\hookrightarrow \text{var}(\beta - \hat{\beta}) = \text{var}(\beta) + \text{var}(\hat{\beta}) - 2\text{cov}(\beta, \hat{\beta}) = \text{var}(\hat{\beta})$

β is fix $\Rightarrow \text{var}(\beta) = 0$
 $\text{cov}(\beta, \hat{\beta}) = 0$

with $\hat{\beta} = (X^T X)^{-1} X^T y$ \rightarrow linear y^T
 quadratic $y^T y$
 y^T

$$\begin{aligned} \text{var}(\hat{\beta}) &= \text{var}[(X^T X)^{-1} X^T y] \\ &= (X^T X)^{-1} X^T \underbrace{\text{var}(y)}_{I \cdot \sigma^2} (X^T X)^{-1} = (X^T X)^{-1} X^T V X (X^T X)^{-1} \\ &= (X^T X)^{-1} \sigma^2 \end{aligned}$$

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- BLUP of u
 $\text{var}(u - \hat{u})$ is called Prediction Error Variance (PEV)
- u is random $\Rightarrow \text{var}(u) = G$
- Compute $\text{var}(u - \hat{u}) = \text{var}(u) + \text{var}(\hat{u}) - 2\text{Cov}(u, \hat{u})$
- Based on properties of BLUP
 $\text{var}(\hat{u}) = \text{cov}(u, \hat{u})$ insert in PEV
where $\hat{u} = G \cdot Z' \cdot V^{-1} (y - X\beta)$
 $\Rightarrow \text{PEV} = \text{var}(u - \hat{u}) = \text{var}(u) + \text{var}(\hat{u}) - 2\text{cov}(u, \hat{u})$
 $= \text{var}(u) - \text{var}(\hat{u})$
- PEV is related to reliability ($B\%$ - Bestimmtheitsmass)
 $B = r_{u, \hat{u}}^2$ with $B\% = 100 \cdot B$

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$$q \text{ PEV}(\hat{u}) = \text{var}(u) - \text{var}(\hat{u}) = C^{22} \quad (4)$$

C^{22} is part of inverse of coefficient Matrix M of mixed model equations

$$\underbrace{\begin{bmatrix} X^T R^{-1} X & X^T R^{-1} Z \\ Z^T R^{-1} X & Z^T R^{-1} Z + G^{-1} \end{bmatrix}}_M \begin{bmatrix} \hat{\beta} \\ \hat{u} \end{bmatrix} = \begin{bmatrix} X^T R^{-1} y \\ Z^T R^{-1} y \end{bmatrix}$$

$\hat{S} = r$

$$M^{-1} = \begin{bmatrix} X^T R^{-1} X & X^T R^{-1} Z \\ Z^T R^{-1} X & Z^T R^{-1} Z + G^{-1} \end{bmatrix}^{-1} = \begin{bmatrix} C^{11} & C^{12} \\ C^{21} & C^{22} \end{bmatrix}$$

Anzahl col $\hat{=}$
Number of animals in pedigree

C^{22} is symmetric with dimensions $q \times q$
with q being the number of animals in pedigree.

OHP Picture 5

□ Reliability of predicted breeding value \hat{u}_i for animal i : (5)

$$B_i = r_{u_i, \hat{u}_i}^2 = \frac{\text{cov}(u_i, \hat{u}_i)^2}{\text{var}(u_i) \text{var}(\hat{u}_i)} = \frac{\text{var}(\hat{u}_i)^2}{\text{var}(u_i) \text{var}(\hat{u}_i)}$$

using BLUP property : $\text{var}(\hat{u}_i) = \text{cov}(u_i, \hat{u}_i)$

$$B_i = \frac{\text{var}(\hat{u}_i)}{\text{var}(u_i)} \Rightarrow \text{var}(\hat{u}_i) = B_i \cdot \text{var}(u_i)$$

$$\text{PEV}(u_i) = \text{var}(u_i) - \text{var}(\hat{u}_i)$$

$$= \text{var}(u_i) - B_i \text{var}(u_i) = (1 - B_i) \text{var}(u_i)$$

↓ solve for B_i

$$B_i \text{var}(u_i) = \text{var}(u_i) - \text{PEV}(u_i)$$

$$B_i = \frac{\text{var}(u_i) - \text{PEV}(u_i)}{\text{var}(u_i)} = 1 - \frac{\text{PEV}(u_i)}{\text{var}(u_i)}$$

$$= 1 - \frac{(G^{22})_{ii}}{\text{var}(u_i)}$$

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□ $MM\bar{E}$

$$\begin{bmatrix} X^T X \\ Z^T X \end{bmatrix} \begin{bmatrix} \hat{\beta} \\ \hat{u} \end{bmatrix} = \begin{bmatrix} X^T y \\ Z^T y \end{bmatrix}$$

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

□ $\begin{bmatrix} X^T X & X^T R^{-1} Z \\ Z^T R^{-1} X & Z^T R^{-1} Z + G^{-1} \end{bmatrix}$ $R^{-1} = I \cdot \sigma_e^{-2}$
 $\underbrace{\hspace{10em}}_{C^{22}}$ $G^{-1} = A^{-1} \cdot \sigma_u^{-2}$

$\lambda = \frac{\sigma_e^2}{\sigma_u^2} = 1$; $\sigma_p^2 = 0.65 = \sigma_e^2 + \sigma_u^2$

$h^2 = \frac{\sigma_u^2}{\sigma_p^2} = \frac{\sigma_u^2}{\sigma_u^2 + \sigma_e^2} = 0.5$ $\sigma_u^2 = \frac{\sigma_p^2}{2} =$

$B_i = 1 - \frac{(C^{22})_{ii}}{\text{Var}(u_i)}$

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□ Meaning:
 individual reliabilities (B_i) make statement about potential risk when using i as parent.

□ Analyse conditional distribution $f(u/\hat{u})$

Eg. animal 3: $\hat{u}_3 = 0.1096$

Normalverteilung
 mit Mean = \hat{u}
 and SD = SEP
 where $SEP = \sqrt{PEV}$

$\alpha/2 = 0.025$

□ Confidence Interval: (95%) $\Rightarrow \alpha = 0.05$

low: $\hat{u}_3 - 1.96 \cdot SEP =$

upper: $\hat{u}_3 + 1.96 \cdot SEP =$

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□ Selection Response (R)

$$R = i \cdot r_{u\hat{u}} \cdot \sigma_u$$

□ For a breeding program:

- R per generation interval = R/L
- 4-path model

$$\frac{R}{L} = \frac{R_1 + R_2 + R_3 + R_4}{L_1 + L_2 + L_3 + L_4} \rightarrow \text{generation interval on each path}$$

$$R_1 = i_1 \cdot r_{u\hat{u}} \cdot \sigma_u$$

$R_2 =$
 $R_3 =$
 $R_4 =$

→ accuracy for all mate selection candidates not individual accuracies.

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□ $r_{u,\hat{u}}$ in selection response

Eg. Path 1:

| | Animal | Ass | Sire | Dam |
|----------------------|--------|---------------------|------|-----|
| Sire | | | | |
| ↓ ① | | | | |
| Sire | | | | |
| Selection candidates | | X delete | | |

Compute $r_{u,\hat{u}}$

① based on correlation between \hat{u} from whole dataset
 \hat{u}_{whole}

② \hat{u}_{reduced} without observation which were deleted

$r_{u,\hat{u}} \Rightarrow r_{\hat{u}_{\text{whole}}, \hat{u}_{\text{reduced}}}$ → Inserted in computation of R