

Evaluation of animals in practice

- Need proper data (centralized)
 - recording system (management groups)
 - correct animal identification
 - other issues?
- Need proper model
 - Account for bias and selection
 - Account for other effects (maternal, permanent environment, multiple trait, different breeds)

A data collection system

- Rules for recording
- Rules for invalid data
- Avoid selective recording
- Rules for defining management groups
- Impetus for doing the right thing
 - What are mechanisms for regulating the system?

Choosing the right model

- Technical issues
 - How to implement the different effects (see next)
- Political issues
 - Evaluation (and reporting) across herds
 - Evaluation across breeds
 - Evaluation across countries
 - Finding a balance between individual and collective satisfaction

More political issues

- Price for the services
- Ownership of the genetic evaluation results
- Ownership of data
-

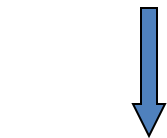
How to expand the simple mixed model

- Simple mixed model

$$y = \text{contempgrp} + \text{animal} + \text{residual}$$

- More general

– $y = \text{fixed effects} + \text{random effects} + \text{residual}$



- cg
- age
-



- animal
- maternal
- permanent env.



homo/
heterogeneous

$$y = \mathbf{Xb} + \mathbf{Zu} + e$$

A sire model

$$\mathbf{y} = \mathbf{Xb} + \mathbf{Zs} + \boldsymbol{\varepsilon}$$

$$\begin{aligned} \text{var}(\mathbf{u}) &= \mathbf{A} \sigma_s^2 \\ \text{var}(\mathbf{e}) &= \mathbf{I} \sigma_\varepsilon^2 \end{aligned} \quad \begin{bmatrix} \mathbf{X}'\mathbf{X} & \mathbf{X}'\mathbf{Z} \\ \mathbf{Z}'\mathbf{X} & \mathbf{Z}'\mathbf{Z} + \lambda \mathbf{A}^{-1} \end{bmatrix} \begin{bmatrix} \mathbf{b} \\ \mathbf{s} \end{bmatrix} = \begin{bmatrix} \mathbf{X}'\mathbf{y} \\ \mathbf{Z}'\mathbf{y} \end{bmatrix}$$
$$\lambda = \sigma_\varepsilon^2 / \sigma_s^2$$

originally used (pre-1985)

fewer equations for amount of data

ignores dam-side

Some formal definitions of the model

$$\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{Z}\mathbf{u} + \mathbf{e}$$

$$\text{var}(\mathbf{u}) = \mathbf{G}$$

$$\text{var}(\mathbf{e}) = \mathbf{R}$$

$$\text{var}(\mathbf{y}) = \mathbf{Z}\mathbf{G}\mathbf{Z}' + \mathbf{R}$$

$$\begin{bmatrix} \mathbf{X}'\mathbf{R}^{-1}\mathbf{X} & \mathbf{X}'\mathbf{R}^{-1}\mathbf{Z} \\ \mathbf{Z}'\mathbf{R}^{-1}\mathbf{X} & \mathbf{Z}'\mathbf{R}^{-1}\mathbf{Z} + \mathbf{G}^{-1} \end{bmatrix} \begin{bmatrix} \mathbf{b} \\ \mathbf{u} \end{bmatrix} = \begin{bmatrix} \mathbf{X}'\mathbf{R}^{-1}\mathbf{y} \\ \mathbf{Z}'\mathbf{R}^{-1}\mathbf{y} \end{bmatrix}$$

simple version

$$\text{var}(\mathbf{u}) = \mathbf{A} \sigma_a^2$$

$$\text{var}(\mathbf{e}) = \mathbf{I} \sigma_e^2$$

$$\begin{bmatrix} \mathbf{X}'\mathbf{X} & \mathbf{X}'\mathbf{Z} \\ \mathbf{Z}'\mathbf{X} & \mathbf{Z}'\mathbf{Z} + \lambda \mathbf{A}^{-1} \end{bmatrix} \begin{bmatrix} \mathbf{b} \\ \mathbf{u} \end{bmatrix} = \begin{bmatrix} \mathbf{X}'\mathbf{y} \\ \mathbf{Z}'\mathbf{y} \end{bmatrix}$$

A simple example of variance structure

animal obs'n

1

9

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

1

11

2

10

$$Z'Z = \begin{bmatrix} 2 & 0 \\ 0 & 1 \end{bmatrix} \quad ZZ' = \begin{bmatrix} 1 & 1 & 0 \\ 1 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

$$G = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \sigma_{anm}^2$$

$$\text{var}(\mathbf{y}) = \mathbf{ZGZ}' + \mathbf{R} = \begin{bmatrix} 1 & 1 & 0 \\ 1 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \sigma_{anm}^2 + \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \sigma_e^2$$

Repeatability model

$$\mathbf{y} = \mathbf{Xb} + \mathbf{Zu} + \mathbf{Zp} + \boldsymbol{\varepsilon}$$

$$\mathbf{G} = \begin{pmatrix} A\sigma_a^2 & 0 \\ 0 & I\sigma_c^2 \end{pmatrix} \quad \text{var} \begin{pmatrix} u \\ p \\ e \end{pmatrix} = \begin{pmatrix} A\sigma_a^2 & 0 & 0 \\ 0 & I\sigma_c^2 & 0 \\ 0 & 0 & I\sigma_e^2 \end{pmatrix} = \begin{pmatrix} \mathbf{G} & 0 \\ 0 & \mathbf{R} \end{pmatrix}$$

$$\begin{pmatrix} X'X & X'Z & X'Z \\ Z'X & Z'Z + \alpha A^{-1} & Z'Z \\ Z'X & Z'Z & Z'Z + \gamma \mathbf{I} \end{pmatrix} \begin{pmatrix} b \\ u \\ p \end{pmatrix} = \begin{pmatrix} X'y \\ Z'y \\ Z'y \end{pmatrix}$$

Maternal effects model

$$y = Xb + Z_1u + Z_2m + \varepsilon$$

Direct animal effect

Maternal effect

$$G = \begin{pmatrix} A\sigma_a^2 & A\sigma_{am} \\ A\sigma_{am} & A\sigma_{mc}^2 \end{pmatrix}$$

Covariance

$$\text{var} \begin{pmatrix} u \\ m \\ e \end{pmatrix} = \begin{pmatrix} A\sigma_a^2 & A\sigma_{am} & 0 \\ A\sigma_{am} & A\sigma_m^2 & 0 \\ 0 & 0 & I\sigma_e^2 \end{pmatrix}$$

$$\begin{pmatrix} X'X & X'Z_1 & X'Z_2 \\ Z_1'X & Z_1'Z_1 + \alpha_{11}A^{-1} & Z_1'Z_2 + \alpha_{12}A^{-1} \\ Z_2'X & Z_2'Z_1 + \alpha_{21}A^{-1} & Z_2'Z_2 + \alpha_{22}A^{-1} \end{pmatrix} \begin{pmatrix} b \\ u \\ m \end{pmatrix} = \begin{pmatrix} X'y \\ Z_1'y \\ Z_2'y \end{pmatrix}$$

Consequence of more complex models

- Usually (many) more equations
- Do we know the parameters (variance components)?
- More difficult to interpret results
- Often more accurate (and less biased)
 - Account for maternal effects
 - Account for heterogeneous variance (animals maybe be more different in some herds/flocks than in other)

Genetic groups

- Consider them as fixed effect in the model
- But add those to breeding values.....
- $EBV_{\text{across}} = EBV_{\text{within}} + \text{group_solution}$
- Grouping needed whenever there is a **genetic** difference in base animals
 - (to account for selection: breeds, origin,....)
- Only need to group the unknown parents
 - Remember that relationships matrix accounts for other selection

Example of genetic groups

Michael Angus	315	Mean Angus	300
Whiskey Hereford	315	Mean Hereford	320

	<u>EBV_{within}</u>	<u>EBV_{across}</u>
Michael Angus	+ 6	+ 6
Whiskey Hereford	- 2	+ 18