How Genomic BLUP Works

Peter von Rohr

09.03.2020

Advantage of Genomic Selection

- Big advantage of genomic selection: shorter generation interval
- Old days: selection mainly based on performance tested sires
 - fine in species where generation inverval is low, e.g. pigs
 - problem when generation interval is long like cattle

Performance Testing



Genomic Selection

- SNP Genotype can be determined when calf is born
- Genomic Breeding value can be computed as soon as SNP genotype is determined
- \blacktriangleright Full sibs get different genomic breeding values \rightarrow increase accuracy

Genomic Selection (II)



Prediction of Genomic Breeding Values

Genomic breeding values can be predicted using either

- Two step approach or
- Single step GBLUP

Two Step Approach



Estimate SNP-effects using a Marker Effect Model

Single Step GBLUP

Use a mixed linear effect model

Genomic breeding values g are random effects

$$y = Xb + Zg + e$$

with

Solution Via Mixed Model Equations

All animals have genotypes and observations

$$\begin{bmatrix} X^T X & X^T Z \\ Z^T X & Z^T Z + \lambda * G^{-1} \end{bmatrix} \begin{bmatrix} \hat{b} \\ \hat{g} \end{bmatrix} = \begin{bmatrix} X^T y \\ Z^T y \end{bmatrix}$$
with $\lambda = \sigma_e^2 / \sigma_g^2$.

Animals Without Observations

- Young animals do not have observations
- Partition ĝ into
 - \hat{g}_1 animals with observations and
 - \hat{g}_2 animals without observations
- Resulting Mixed Model Equations are (assume $\lambda = 1$)

$$\begin{bmatrix} X^T X & X^T Z & 0 \\ Z^T X & Z^T Z + G^{(11)} & G^{(12)} \\ 0 & G^{(21)} & G^{(22)} \end{bmatrix} \begin{bmatrix} \hat{b} \\ \hat{g}_1 \\ \hat{g}_2 \end{bmatrix} = \begin{bmatrix} X^T y \\ Z^T y \\ 0 \end{bmatrix}$$

Predicted Genomic Breeding Values

$$G^{(21)} \cdot \hat{g}_1 + G^{(22)} \cdot \hat{g}_2 = 0$$

Solutions



$$\hat{g}_2 = -(G^{(22)})^{-1} \cdot G^{(21)} \cdot \hat{g}_1$$