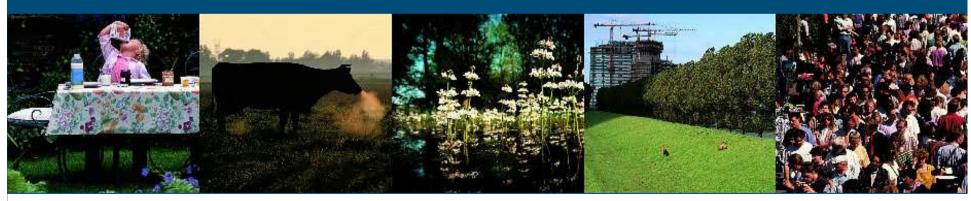
Accuracies of EBVs:

Standard Errors of EBVs vs. Response to Selection

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The role of accuracy (ρ) in animal breeding

- 1. Response to selection: $R = \frac{i\rho\sigma_A}{L}$
 - ρ = correlation(EBV,BV)
 - ρ measures the quality of a breeding scheme

- 2. Credibility of an EBV: $SE(EBV) = \sqrt{(1-\rho^2)\sigma_A^2}$
 - ρ measures the quality of a single EBV
 - Risk that EBV changes with more info
 - Dairy cattle, horses

Accuracies are routinely produced in BVE

- BVE → Accuracies follow from the PEV: $\rho_{PEV} = \sqrt{\frac{\sigma_A^2 PEV}{\sigma_A^2}}$
 - Inverse of coefficient matrix of MME
- ρ_{PEV} is a convenient proxy for response to selection
 - Often routinely available
 - Easily compare individuals with different info
- Example: Dairy cattle
 - Parent Average: $\rho_{PEV} = 0.55$ (rel. = 0.3)
 - + 1st lactation: $\rho_{PFV} = 0.64 (+15\%)$

Conclusion:

Waiting for 1st lactation record

yields ~15% more response

- However, ρ_{PFV} ignores selection (Selection matters: Dekkers, 1992 Anim. Prod. 54)
 - The above may be somewhat erroneous

Objective

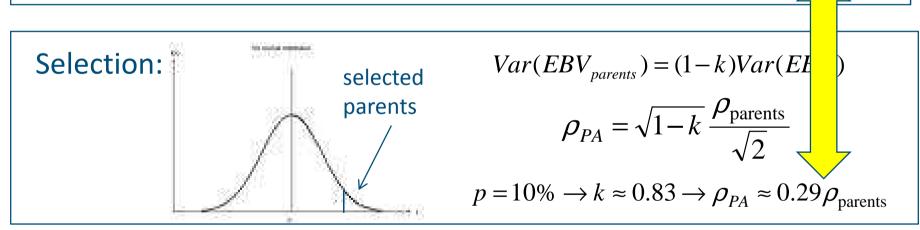
- To demonstrate that:
 - Ordinary accuracies of EBVs may <u>not at all</u> reflect response to selection
 - The error can be much greater than the usual Bulmer-effect
 - SE(EBV) and Corr(EBV,BV) are different things
 - There exists no single "accuracy" reflecting both measures

The accuracy of the Parent Average (PA)

Accuracy defined as correlation(EBV,BV)

$$PA = (EBV_S + EBV_D)/2$$

No selection:
$$Var(PA) = \frac{1}{2} Var(EBV_{parents}) \rightarrow \rho_{PA} = \frac{\rho_{parents}}{\sqrt{2}} \approx 0.71 \rho_{parents}$$



Selection strongly reduces $Var(EBV) \rightarrow \rho_{PA}$ is strongly reduced

Ordinary accuracy from PEV ignores selection

No selection:
$$\sigma_{EBV}^2 = \sigma_A^2 - PEV \rightarrow \rho_{PEV} = \sqrt{\frac{\sigma_A^2 - PEV}{\sigma_A^2}} = \sqrt{\frac{\sigma_{EBV}^2}{\sigma_A^2}} = corr(A, EBV)$$

Selection:
$$\sigma_{EBV}^2 \neq \sigma_A^2 - PEV \rightarrow \rho_{PEV} = \sqrt{\frac{\sigma_A^2 - PEV}{\sigma_A^2}} \neq \sqrt{\frac{\sigma_{EBV}^2}{\sigma_A^2}} = corr(A, EBV)$$

With selection,
the standard error of an EBV (VPEV)
and the correlation between true and estimated BV
are different things.

Comparison of breeding schemes

- The ordinary accuracy systematically overpredicts the value of pedigree information
- This biases the comparison of breeding schemes
 - New information is undervalued
 - Own performance
 - Genomic information
 - Progeny

Example: value of genotyping cows in GS

- Milk yield
 - GS-accuracy = 0.8
 - Selected proportions: 1% in both sexes
 - Bulls are genotyped
- Issue: The effect of genotyping cows on accuracy of their EBV

| Method | Parent Avg. | Genotyping | Difference |
|--------------|-------------|------------|------------|
| ρ_{PEV} | 0.43 | 0.78 | +81% |
| Corr(EBV,BV) | 0.12 | 0.68 | +466% |

5-fold difference

Ordinary accuracy undervalues genomic info in cows

Example: GS for litter size in pigs

- Litter size
 - GS-accuracy = 0.4
 - Selected proportions: 25% in sows, ~2% in boars
 - Selection sows after 1st litter.
 - Info: 1st litter, 3FS, 56HS records
 - Selection of boars: at sexual maturity
 - Info: Parent Average
- Issue: Select boars on Parent Average or Genomic Selection?

| Method | Parent Avg. | Genotyping | Difference |
|--------------|-------------|------------|------------|
| ρ_{PEV} | 0.31 | 0.52 | +66% |
| Corr(A,EBV) | 0.13 | 0.36 | +182% |

2.75-fold difference

Conclusions

- Ordinary accuracies obtained from BVE:
 - Reflect the standard error of an EBV
 - Do not reflect response to selection
 - Cannot be used to compare breeding schemes
 - Overpredict the value of pedigree information for response
- Accuracies relevant for response can be obtained from:
 - Selection-index calculations
 - e.g. SelAction, not SELIND
 - SD_{FBV} /SD_{BV} for Total Merit in the selection candidates
 - (Cross)validation using historical data
- Dairy cattle breeders mix two incompatible concepts of accuracy
 - Monday's session

