Integration of external information from multiple traits into the national multitrait evaluations



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Background and Objective

Inclusion of Interbull MACE information to national single-step genomic evaluations (blending) is crucial

Steps for blending

- 1. Calculate multi-trait effective record contribution
- Frequently MACE is combined trait, national evaluations multi-trait models
- We tested the efficiency of blending combined indices (protein, milk, fat) to multi-trait multi-parity model

Data and Models

The blending method was studied using 305d milk, protein, and fat yields for first three lactations for Nordic (FIN, SWE, DNK) Holstein cows

- Three multi-trait evaluation models
 - 1. Domestic evaluation using **Danish data** (**DNK**)
 - 2. External Nordic evaluation using **Finnish**,
 - **Danish**, and **Swedish** data (**DFS**)

- (ERC) and multi-trait deregressed proofs (DRP) for domestic and external trait indices
- 2. Amount of external information

 $ERC_{bl} = ERC_{ext} - ERC_{dom}$

- 3. Pseudo observation $DRP_{bl} = \frac{ERC_{ext} * DRP_{ext} - ERC_{dom} * DRP_{dom}}{ERC_{ext} - ERC_{dom}}$
- 4. Include DRP_{bl} to domestic model as an observation using *ERC_{bl}* as a weight





- 3. Domestic evaluation model including external information
- 9 traits = 3 lactations, 3 traits within lactation
- **Genetic correlation** between countries = 1
- Trait indices were calculated based on breeding \bullet values for parities as $0.5 * EBV_1 + 0.3 * EBV_2 + 0.2 EBV_3$
- A **bull** was blended if
 - 1. Reliability of protein index in DFS was >0.5 and was **0.05 units higher** compared to **DNK** 2. Birth year of a bull was >1989

In total **11738** bulls were selected to be blended

Domestic

External information

Blended



Conclusions

- Blending of external information as trait indices to a multi-trait multi-parity evaluation model works
- The regression slope and correlation between blended and DFS protein indices are about 1
- No bias in reliability

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