# Improving the model for genetic evaluation of calving traits in the US Holstein and Brown Swiss



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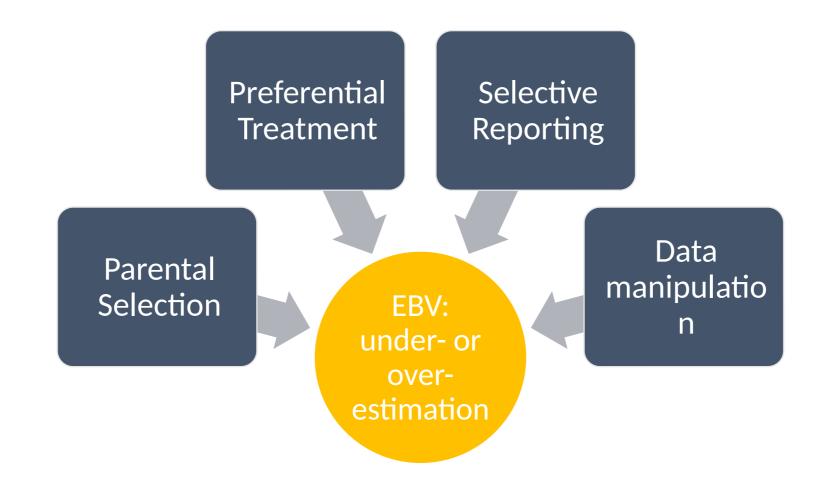
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# Background



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2. an additional and possibly underestimated source of bias: excluding one source of variation from the model

- preadjusting milk production for age and parity under- or over-estimation of the genetic trend
- Lidauer and Mäntysaari (1996): redefining the herd effect in the Finnish repeatability animal model evaluation bias
- ICAR's guidelines on Dairy Cattle Genetic Evaluation : the importance of model's unbiasedness

## Aim

The aim of the present study was to

- 1. Investigate possible improvements of the current National Genetic evaluation for Calving Ease (CE) and Stillbirth (SB) in the US Holstein and Brown Swiss
- 2. Validate results by ITB methods 1 and 3

## Data

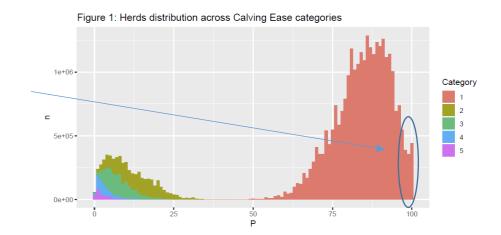
- CDCB Data from the official April 2019 run were used
  - Calving Ease (CE): 32,194,410 records
    - 1st parity 29.8%, 2nd parity 29.3%, 3+ parities 40.9 %
  - Stillbirth (SB): 20,902,357 records
    - 1st parity 30.9%, 2nd parity 29.2%, 3+ parities 39.9 %

### Current Model

- Current model for calving traits (Van Tassell et al, 2003; Cole et al, 2005)
  - Single-trait threshold sire-maternal grandsire (MGS) model
  - environmental effects: Random herd-year, fixed year-season, parity-sex, sire/ mgs birth year group, MGS breed (CE only)
  - PTA: % difficult births (CE score 4 & 5) in heifers and percent stillbirths (score 2 & 3) over all parities

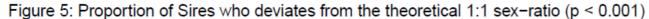
# Preliminary Analyses

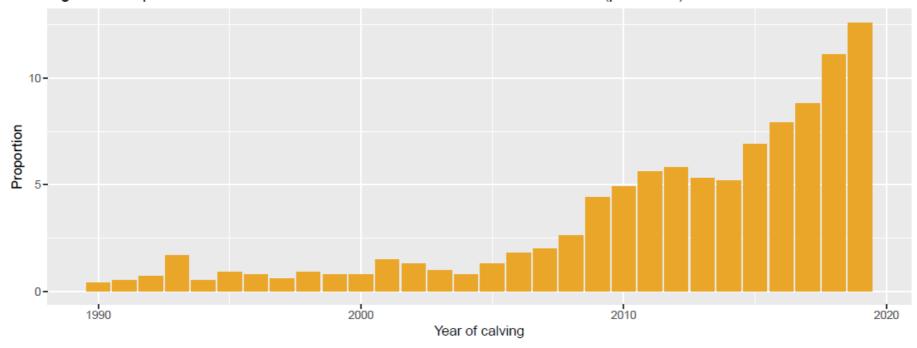
- Possible model improvements
  - Exclude herds with > 95% of easy calvings/no stillbirth
  - 2. CE: join category 4 & 5
  - 3. Include parity in the definition of HY groups
  - 4. Include the interaction of Parity-Sex-Year of birth of Sire and MGS



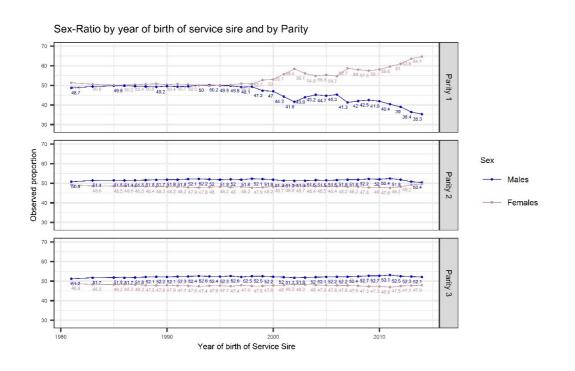
## Has the sex-ratio by sire changed across years?

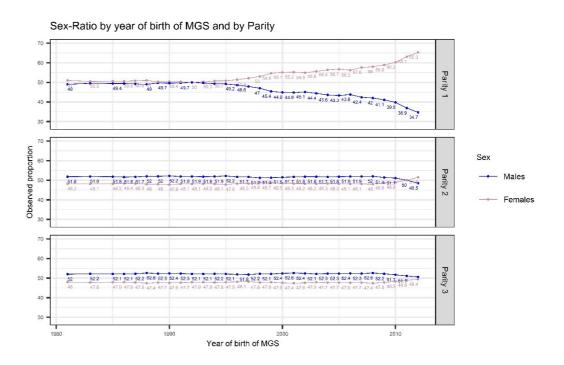
#### YES!





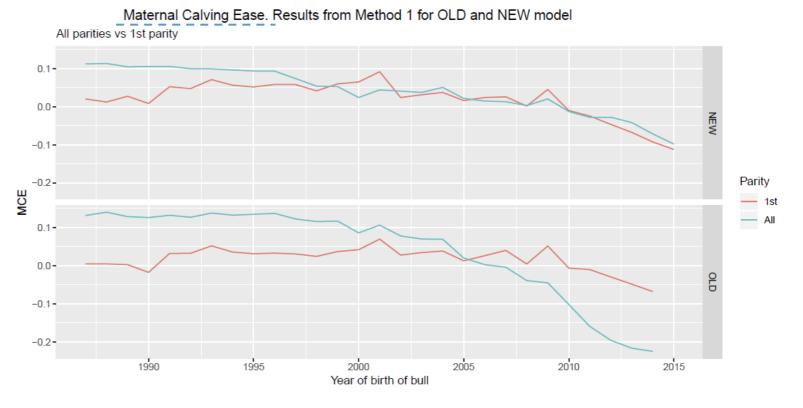
#### Has the sex-ratio by sire/MGS changed across years?





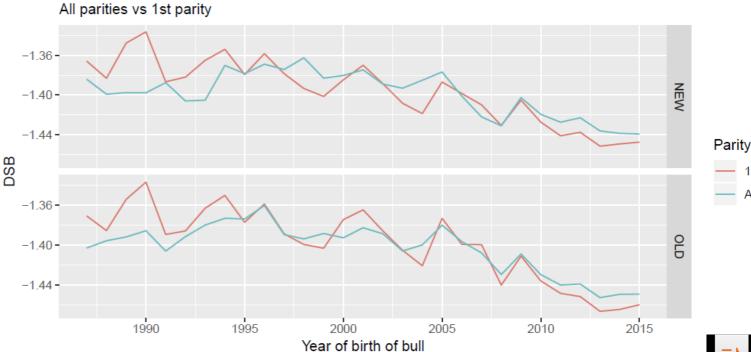
## Results – ITB method 1

- routine national genetic evaluations: compares genetic trends estimated using only first lactation versus all lactations
  - All traits passed ITB 1
  - Largest impact on CE



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Direct Stillbirth. Results from Method 1 for OLD and NEW model



## Results – ITB method 3

 routine national genetic evaluations: analyses the official national predicted genetic merit variation across evaluation runs

• All traits passed ITB 3 test

Largest impact on CE

Method 3 - Comparison between OLD and NEW model 0.05 -0.00 --0.05 -Direct CE PTA RUN 0.05 -0.00 --0.05 -2008 2016 2004 2012 Sire year of birth

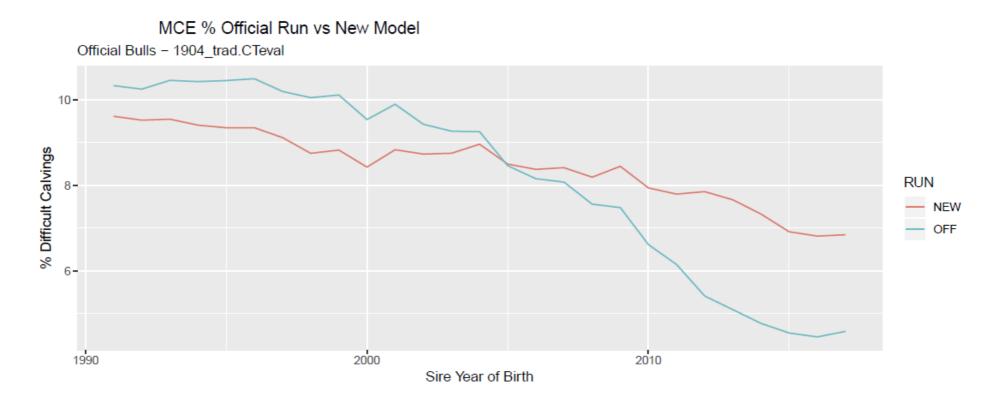
## Results - Genetic Correlation

- The introduced changes increased the international genetic correlation between US and 14, 7, 1 and 2 countries for DCE, MCE, DSB and MSB, respectively.
- The largest observed effects were actually for both DCE and MCE
- The new variance components and the new model have had an effect on the reliabilities of US bulls across different countries.



## Results - Genetic Trend

model improvement: effect on the genetic trends (especially for MCE)





## Conclusion and Final remarks

- evidence of a bias due to sexed-semen
- new model: Inclusion of the interaction of Parity-Sex-Year of birth of Sire and MGS.
- positive results in terms of validation tests
- positive results in terms of genetic correlation with the other countries (on average)
- effect on genetic trends (MCE)



## Thank you for the attention

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