Session 39. Breeding for robustness in cattle – part 2. Presentation no. 2
Book of abstracts page 382

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Genetic Analysis of Stillbirth and Calving Difficulty in Norwegian Red cows Bjørg Heringstad¹², Yu-Mei Chang³, Morten Svendsen¹, and Daniel Gianola².³ ¹Geno Breeding and A. I. Association, Ås, NORWAY ²Dep of Animal and Aquacultural Sci, Norwegian University of Life Sciences, Ås, NORWAY ³ Department of Dairy Science, University of Wisconsin, Madison, USA 58th Annual meeting of the EAAP, Dublin, Irland, August 26-29, 2007 Commission on Animal Genetics, Session: Breeding for robustness in cattle, 39:2

Objectives



- Infer genetic parameters for calving difficulty and stillbirth using a bivariate sire-maternal grandsire threshold model
- Evaluate phenotypic and genetic change for these traits for Norwegian Red cows

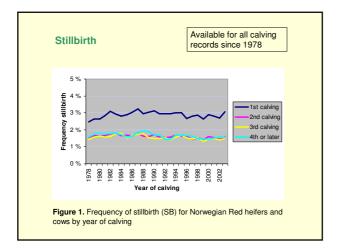
Traits

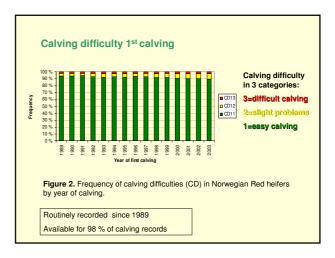
Stillbirth is a binary trait:

- 0 = born alive
- 1 = dead at birth or within 24 hours

Calving difficulty in 3 categories:

- 1 = easy calving
- 2 = slight problems
- 3 = difficult calving





Calving difficulty 2nd and later calving

CD score similar for all parities after 1st calving Overall mean:

- 97 % easy calving
- 2 % slight problems
- 1 % difficult calving

Data

ion calving records

- 528,475 first-lactation calving records
- First calving 1989-2004
- Age at first calving 21-34 mo
- From herd-year classes with at least 5 first lactation cows
- Known Norwegian Red A.I. sire as sire of calf and sire of cow (maternal grandsire)
 - 1548 Norwegian Red sires with records
 - 2155 in pedigree-file

Data

Distribution of records over SB categories

SB		No of records	Frequency
0	born alive	514,393	97.3 %
1	dead at birth	14,082	2.7 %

Distribution of records over CD categories

CD		No of records	Frequency
1	Easy calving	475,945	90.1 %
2	Slight problems	27,142	5.1 %
3	Difficult calving	12,684	2.4 %
4	Unknown	12,704	2.4 %



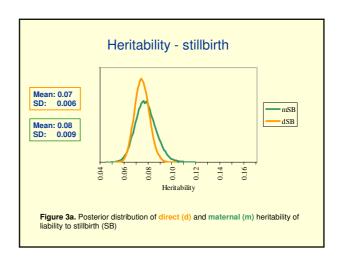
Model

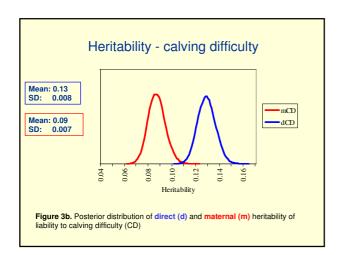
- Bivariate sire maternal grandsire thresholdliability model for one binary (SB) and one ordered categorical (CD) trait
- Bayesian approach using MCMC methods

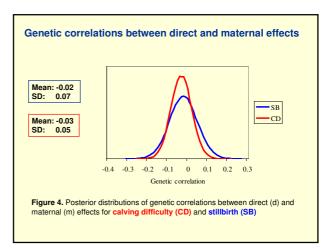
Bivariate sire - mgs threshold-liability model

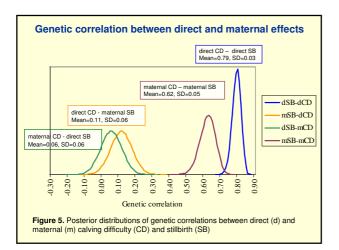
$$\lambda = X\beta + Z_{_h}h + Z_{_u}u + e$$

- λ vector of unobserved liabilities of SB and CD
- β vector of systematic effects sex of calf (2 levels), age (14 classes) and mo-yr (168 classes) of first calving
- h vector of herd-year of calving effects (85,255 levels)
- $\begin{array}{ll} \textbf{u} & \text{vector of effects of the sire (s) and maternal grandsire} \\ & (\text{mgs}) \text{ of the calf} & \textbf{u}' = [\textbf{s}'_{\textit{cD}} \textbf{mgs}'_{\textit{cD}} \textbf{s}'_{\textit{sB}} \textbf{mgs}'_{\textit{sB}}] \end{array}$
- e vector of residual effects
- $\boldsymbol{X},\,\boldsymbol{Z}_h,$ and \boldsymbol{Z}_u incidence matrices



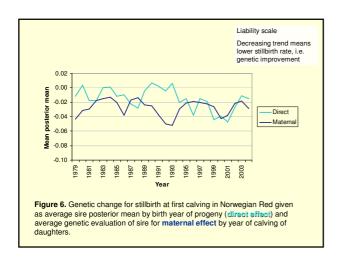


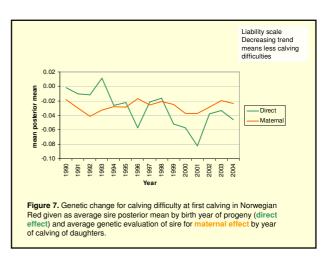




Genetic change

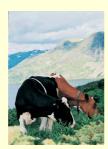
- Average sire posterior means by birth year of progeny (direct) or year of calving of daughters (maternal).
- All 1st calving records of the 1548 sires/mgs (900.000 records) were used for assessment of genetic change.
- Sires were weighted according to their number of daughters, so this measure reflects sire usage as well as possible genetic change in the Norwegian Red population.





Summary and conclusions

- Low levels of stillbirth and calving difficulty in the Norwegian Red population (2% SB, 95% easy calving)
- No genetic change for SB and a slight genetic improvement for CD were found in the Norwegian Red population.



Heringstad, B., Y. M. Chang, M. Svendsen, and D. Gianola. 2007. Genetic analysis of calving difficulty and stillbirth in Norwegian Red cows. Journal of Dairy Science 90: 3500-3507.

