

Association of genomic and parental breeding values with cow performance in Nordic dairy cattle

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AIM

Compare genomically enhanced

BACKGROUND

Genotyping females in the

METHODS

Data were collected from the

breeding values (GEBV) and parent average breeding values (PA) for heifers regarding their ability to predict the cow's future performance.

CONCLUSION

Genomically enhanced breeding values predicted cow performance significantly better than parent average breeding values for the vast majority of the analyzed traits. Nordic countries started on a large scale in 2012. Since the start, over 250,000 females have been genotyped.

 Validation of GEBV and illustrating the relationship between genomic prediction and the future phenotype is key to increasing confidence in the genomic technology among farmers.

Swedish, Danish and Finnish milk recording schemes.

- Production, conformation, fertility and other functional traits, adjusted for systematic environmental effects, were used as measures of cow performance.
- GEBV and PA were from the Nordic Cattle Genetic Evaluation.

RESULTS

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| Cow | Breeding | Correlation | Correlation |
|-------------|----------|-------------|-------------|
| Performance | Value | with GEBV* | with PA* |

0.36



0.25

Protein Index, Protein_{Adi} Red Dairy Cattle

18

| | SCS _{Adj} | Mastitis | -0.21 | -0.13 |
|----------|------------------------|-----------|-------|-------|
| Holstein | IFL _{Adj} | Fertility | -0.11 | -0.07 |
| | | | | |
| | Protein _{Adj} | Protein | 0.40 | 0.24 |
| | SCS _{Adj} | Mastitis | -0.18 | -0.11 |
| Jersey | IFL _{Adj} | Fertility | -0.08 | -0.04 |
| | | | | |
| | Protein _{Adj} | Protein | 0.31 | 0.21 |
| | SCS _{Adj} | Mastitis | -0.16 | -0.10 |
| RDC | IFL _{Adj} | Fertility | -0.09 | -0.05 |

*Negative correlations are desirable for mastitis and fertility





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