

News genomic prediction - NAV routine evaluation 2 June 2014

The latest NAV routine genomic prediction took place as scheduled. NAV carried out genomic prediction for Holstein, RDC and Jersey:

Data used in genomic prediction

Genotypes were extracted from the joint Nordic SNP data base 21th May 2014. Interbull information was from April 2014. Nordic phenotypic information for the reference population used in the genomic prediction was the same as in the 2nd May 2014 routine evaluation.

General about Nordic Cattle Genetic Evaluation: www.nordicebv.info

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