

News genomic prediction - NAV routine evaluation 2 December 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 19th November 2014. Interbull information was from August 2014. Nordic phenotypic information for the reference population used in the genomic prediction was the same as in the 2nd November 2014 routine evaluation.

News in relation to NAV genomic prediction

No changes.

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

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