

News genomic prediction - NAV routine evaluation 2 January 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 15th December 2013. The annual exchanges of genotypes within Eurogenomics for Holstein have taken place and genotypes from the last birth year of bulls with milking daughters are included in the reference population. Interbull information was from December 2013. Nordic phenotypic information used in the genomic prediction was the same as in the 2nd November 2013 routine evaluation.

News in relation to NAV genomic prediction

No changes

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

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