

News genomic prediction - NAV routine evaluation 7 April 2015

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 18th March 2015. Nordic phenotypic information for the reference population used in the genomic prediction was the same as in the 3rd February 2015 routine evaluation.

News in relation to NAV genomic prediction

No changes

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

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