

News genomic prediction - NAV routine evaluation

2 June 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 11th May 2015. INTERBULL information from April 2015, and national information from 5th May run were included in the genomic prediction.

NAV has discovered that the majority of the 88 animals with accepted mismatch corrections since last evaluation did not get GEBVs 2nd of June 2015. NAV has identified what is causing the problem, and the animals will be included in the next publication 7th of July.

News in relation to NAV genomic prediction

In the May run genotyped cows having fertility phenotypes included in the fertility EBV got an EBV published. In the June run genotyped cows get the GEBV published for fertility.

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

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