

News genomic prediction - NAV routine evaluation

15 September 2012

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 28nd August 2012. Interbull information from August 2012. Nordic phenotypic information used in the genomic prediction was the same as in the 14th August 2012 routine evaluation.

News in relation to NAV genomic prediction

Germany participated in Interbull evaluation for calving traits for the first time. It means significantly more information has been included in Holstein reference population about calving traits. Compared to August run an error has been corrected in the RDC GEBVs for claw health. August GEBVs for claw health will be corrected

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