News genomic prediction - NAV routine evaluation 2 October 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 September 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 August 2014 routine evaluation.

Changes

For RDC we observed some instability for temperament caused by cow DRPs. To fix that problem cows excluded from the RDC reference population for temperament in the October 2nd 2014 run. NAV has to study in details what is causing the problem.

General about Nordic Cattle Genetic Evaluation: www.nordicebv.info
Contact person: Gert Pedersen Aamand, Ph.: +45 87405288 gap@vfl.dk,

Denmark: https://www.landbrugsinfo.dk

Contact person: Ulrik Sander Nielsen, Danish Cattle, Ph. +45 87405289, usn@vfl.dk

Sweden: www.sweebv.info, www.vxa.se

Contact person: Jan-Åke Eriksson, Swedish Dairy Association, Ph. +46 08-790 58 67

Jan-Ake.Eriksson@vxa.se

Finland: www.faba.fi

Contact person: Jukka Pösö, Faba co-op, Ph +358-(0)207472071 jukka.poso@faba.fi