

## ***News genomic prediction - NAV routine evaluation 6 October 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 21<sup>th</sup> September 2015. INTERBULL information from August 2015 and national information from 11<sup>th</sup> August run were included in the genomic prediction.

### **Genomic prediction for type**

NAV has found that some genotyped Holstein cows having a phenotype have been missing in the reference population. These cows are added to the reference population and their GEBVs include now both the genomic information and the phenotypic information. Earlier it was only the genomic information.

**General about Nordic Cattle Genetic Evaluation:** [www.nordicebv.info](http://www.nordicebv.info)

Contact person: Gert Pedersen Aamand, Ph.: +45 87405288 [gap@seges.dk](mailto:gap@seges.dk),

**Denmark:** <https://www.landbrugsinfo.dk>

Contact person: Ulrik Sander Nielsen, SEGES, Cattle, Ph. +45 87405289, [usn@seges.dk](mailto:usn@seges.dk)

**Sweden:** [www.sweebv.info](http://www.sweebv.info), [www.vxa.se](http://www.vxa.se)

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

[Jan-Ake.Eriksson@vxa.se](mailto:Jan-Ake.Eriksson@vxa.se)

**Finland:** [www.faba.fi](http://www.faba.fi)

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