

## INTRODUCTION

The latest genomic test international evaluation for longevity trait took place as scheduled at the Interbull Centre. Data from 21 populations were included in this evaluation.

International genetic evaluations for direct longevity of bulls were computed from: AUS BEL CAN CHE CZE DEU DFS ESP FRA GBR HUN IRL ISR ITA NLD NZL POL SVN USA ZAF JPN Holstein breed data were included in this evaluation.

CAN, DEU, ESP, FRA, DFS, GBR, ITA, NLD, HUN, POL submitted GEBVs.

dlo: CAN, DEU, ESP, FRA, DFS, GBR, ITA, NLD, HUN, POL

## CHANGES IN NATIONAL PROCEDURES

Changes in the national genetic evaluation of longevity traits are as follows:

FRA (HOL) Proofs and reliability calculated with the single step methodology (HSSGBLUP) developed by INRAE. A new software for the count of daughters and herds has also been developed by Geneval. Principles stayed the same than before but pedigree corrections have been made. Other information concerning publication can have been changed

GBR (HOL) Updates in genotypes and data update

## INTERBULL CHANGES COMPARED TO THE DECEMBER ROUTINE RUN

No changes in Interbull procedures

## DATA AND METHOD OF ANALYSIS

Thirteen Holstein populations sent GEBV data for up to 38 traits, while classical EBVs for the same traits were used in the analyses. Young bull GEBVs from the GEBV providers have been converted to the scales of all countries participating in classical MACE. A bull will get a MACE EBV or a GMACE EBV but not both.

From those thirteen countries, National GEBVs of bulls less than seven years of age and with no classical MACE proofs were included for the breeding value prediction with a further requirement of either a MACE-PA or a GMACE-PA (for young genomic bulls with young genomic sires) being available.

The parameter-space approach is used for the GMACE genetic evaluations (Sullivan, 2016)

## SCIENTIFIC LITERATURE

The international genetic evaluation procedure is based on international work described in the following scientific publications:

Sullivan, P.G. 2016. Defining a Parameter Space for GMACE. Interbull Bulletin 50, p 85-93.

VanRaden, P.M. and Sullivan, P.G. 2010. International genomic evaluation methods for dairy cattle. Gen. Sel. Evol. 42:7

Sullivan, P.G. and Jakobsen, J.H. 2012. Robust GMACE for young bulls methodology. Interbull Bulletin 45, Article 1.

Sullivan, P.G. 2012a. GMACE reliability approximation. Report to the GMACE working group of Interbull. GMACE\_rels 2013

Sullivan, P.G. 2012b. GMACE variance estimation. Report to the GMACE working group of Interbull. GMACE\_vce 2013

Sullivan, P.G. 2012c. GMACE Weighting Factors. Report to the GMACE working group of Interbull. GMACE\_gedcs 2013

Jakobsen, J.H. and Sullivan, P.G. 2013. Trait specific computation of shared reference population. Reference sharing Nov 2013

## NEXT ROUTINE INTERNATIONAL EVALUATION

Dates for next routine run can be found on <http://www.interbull.org/ib/servicecalendar>

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NEXT TEST INTERNATIONAL EVALUATION  
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Dates for next test run can be found on <http://www.interbull.org/ib/servicecalendar>

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PUBLICATION OF INTERBULL ROUTINE RUN  
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Results were distributed by the Interbull Centre to designated representatives in each country. The international evaluation file comprised international proofs expressed on the base and unit of each country included in the analysis. Such records readily provide more information on bull performance in various countries, thereby minimising the need to resort to conversions.

At the same time, all recipients of Interbull results are expected to honour the agreed code of practice, decided by the Interbull Steering Committee, and only publish international evaluations on their own country scale. Evaluations expressed on another country scale are confidential and may only be used internally for research and review purposes.

Table 1. National evaluation dates in GMACE run December 2022

Country	Date
CAN	20221201
DEU	20221206
DFS	20221101
ESP	20221115
FRA	20221206
GBR	20221109
ITA	20221111
NLD	20221201
HUN	20211122
POL	20221110

Table 2.

Number of bulls in reference population for	dlo
CAN 43075.0	
DEU 10058.0 45647.0	
DFS 5588.0 38871.0 39799.0	
ESP 6681.0 40234.0 38986.0 41287.0	
FRA 4154.0 34937.0 34408.0 35029.0 36714.0	
GBR 36372.0 10607.0 5978.0 7165.0 4207.0 38596.0	
ITA 37129.0 9533.0 4980.0 6079.0 3378.0 36318.0 38507.0	
NLD 4221.0 36802.0 36147.0 36772.0 34416.0 4555.0 3552.0 38676.0	
HUN 2274.0 8255.0 7682.0 8077.0 7296.0 2494.0 2256.0 7827.0 9098.0	
POL 4895.0 33671.0 33349.0 33982.0 30468.0 5221.0 4281.0 31969.0 7642.0 35379.0	