INTRODUCTION

The latest genomic routine 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 trait of bulls from Australia, Belgium, Canada, Switzerland, Germany, Denmark-Finland-Sweden Spain, France, The United Kingdom, Ireland, Israel, Italy, New Zealand, The Netherlands, The United States of America Hungary, Norway, Slovenia and Czech Republic were computed. Holstein breed data were included in this evaluation.

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

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

CHANGES IN NATIONAL PROCEDURES

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

DEU (HOL) Base change
CAN (HOL) Base change
ITA (HOL) Base change
GBR (HOL) Base change
FRA (HOL) Base change
NLD (HOL) Base change
POL (HOL) Base change

INTERBULL CHANGES COMPARED TO THE DECEMBER ROUTINE RUN

No changes in Interbull procedures

DATA AND METHOD OF ANALYSIS

Eleven 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 eleven 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

NEXT TEST INTERNATIONAL EVALUATION

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

PUBLICATION OF INTERBULL ROUTINE RUN

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 April 2020

Country	Date
BEL	20190901
CAN	20200401
DEU	20200407
DFS	20200203
ESP	20200317
FRA	20200408
GBR	20200225
ITA	20200309
NLD	20200401
HUN	20200323

Table 2.

Number of bulls in reference population for	dlo								
BEL	3220.0								
CAN	1780.0	36392.0							
DEU	1323.0	5957.0	40204.0						
DFS	1101.0	3830.0	36120.0	36949.0					
ESP	1250.0	4204.0	36736.0	36237.0	37549.0				
FRA	1286.0	3897.0	34298.0	33838.0	34384.0	36243.0			
GBR	1355.0	32887.0	5999.0	3862.0	4234.0	3905.0	34644.0		
ITA	1631.0	32405.0	4806.0	2980.0	3259.0	2997.0	31101.0	32948.0	
NLD	1234.0	3869.0	35747.0	35301.0	35805.0	34004.0	3936.0	2922.0	37807.0
HUN	778.0	1716.0	7322.0	6948.0	7183.0	6933.0	1728.0	1547.0	7172.0	7868.0