

## INTRODUCTION

The latest genomic international evaluation for dairy production traits took place as scheduled at the Interbull Centre. Data 32 countries were included in this evaluation.

International genetic evaluations for milk, fat and protein yields of bulls from Australia, Austria-Germany, Belgium, Canada, Czech Republic, Denmark-Finland-Sweden, Estonia, France, Hungary, Ireland, Israel, Italy, Japan, Latvia, Lithuania, Netherlands, New Zealand, Norway, Poland, Republic of South Africa, Slovak Republic, Slovenia, Spain, Switzerland, the United Kingdom, the United States of America, Portugal, Korea, Argentina and Uruguay were computed. Holstein breed data were included in this evaluation.

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

fat: BEL, CAN, DEU, ESP, FRA, AUS, DFS, GBR, ITA, NLD, POL, HUN  
mil: BEL, CAN, DEU, ESP, FRA, AUS, DFS, GBR, ITA, NLD, POL, HUN  
pro: BEL, CAN, DEU, ESP, FRA, AUS, DFS, GBR, ITA, NLD, POL, HUN

## CHANGES IN NATIONAL PROCEDURES

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

ESP (HOL) Base change

NLD (HOL) Included a deregression post-processing step to keep the animals with information in the system

BEL (HOL) Corrected a small bug in their routines preparing final GEBV to be submitted  
Increase in the size of the reference population (mainly females)

## 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



ITA	30349.0	2890.0	2214.0	2575.0	30938.0								
NLD	3012.0	33789.0	33642.0	32422.0	2388.0	35974.0							
GBR	30507.0	3324.0	2725.0	3171.0	29067.0	2937.0	31727.0						
AUS	1313.0	729.0	624.0	711.0	865.0	737.0	1460.0	4389.0					
BEL	1460.0	1062.0	861.0	1084.0	1311.0	999.0	1020.0	365.0	2798.0				
ESP	3187.0	34355.0	34352.0	33247.0	2493.0	34147.0	3122.0	695.0	1015.0	35937.0			
HUN	1104.0	6231.0	5948.0	6077.0	1013.0	6190.0	1076.0	455.0	494.0	6206.0	6780.0		
POL	3339.0	29486.0	29633.0	28759.0	2643.0	29540.0	2923.0	645.0	1527.0	30393.0	6199.0	32014.0	

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Number of bulls in reference population for                    pro  
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CAN	36769.0												
DEU	3486.0	36577.0											
DFS	2795.0	33819.0	35043.0										
FRA	3269.0	32488.0	32340.0	35107.0									
ITA	30349.0	2890.0	2214.0	2575.0	30937.0								
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