#### INTRODUCTION

The latest genomic test international evaluation for udder traits took place as scheduled at the Interbull Centre. Data from 21 countries were included in this evaluation.

International genetic evaluations for udder health traits 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 data were included in this evaluation.

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

mas: , CAN, DEU, ESP, FRA, DFS, , ITA, NLD, POL, scs: BEL, CAN, DEU, ESP, FRA, DFS, GBR, ITA, NLD, POL, HUN

CHANGES IN NATIONAL PROCEDURES

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

FRA (HOL) Decrease in reliability for mas for a large number of bulls as the parameters to compute reliabilities have been reworked, causing an important downward reliability variation

Changes in information due to pedigree verification

DEU (HOL) Base change

GBR (HOL) Change in bulls' status due to re-assiging codes to bulls as more information from daughters become more available

HUN (HOL) For mas data from 2312r used due to the mismatch between MACE and GMACE parameter file

### INTERBULL CHANGES COMPARED TO THE DECEMBER ROUTINE RUN

No changes in Interbull procedures

### DATA AND METHOD OF ANALYSIS

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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

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

### NEXT TEST INTERNATIONAL EVALUATION

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

# PUBLICATION OF INTERBULL ROUTINE RUN

Country Date

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 2024

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NLD 3598.0 24366.0 24095.0 23391.0 25753.0

20241201 DEU 20241203 DFS 20241105 FRA 20241204 GBR 20241111 NLD 20241201 20241101 ITA HUN 20241115 BEL 20201201 ESP 20241119 20240930 Table 2. Number of bulls in reference population for scs \_\_\_\_\_ CAN 45930.0 DEU 12584.0 49311.0 DFS 6837.0 40970.0 42024.0 FRA 5770.0 37858.0 37253.0 39715.0 GBR 38924.0 13646.0 7750.0 6235.0 41924.0 NLD 4249.0 36918.0 36387.0 34930.0 4576.0 38786.0 ITA 40047.0 12621.0 6677.0 5268.0 39419.0 3601.0 42071.0 HUN 2288.0 8279.0 7803.0 7642.0 2508.0 7827.0 2273.0 9113.0 BEL 729.0 728.0 652.0 719.0 686.0 741.0 722.0 549.0 1719.0 ESP 7841.0 42326.0 40568.0 37945.0 8809.0 36876.0 7674.0 8097.0 705.0 43406.0 POL 5550.0 35059.0 35021.0 32989.0 6285.0 32028.0 5350.0 7640.0 994.0 35264.0 36852.0 \_\_\_\_\_ Number of bulls in reference population for mas \_\_\_\_\_ CAN 27432.0 DEU 10191.0 33786.0 DFS 5742.0 27986.0 28837.0 FRA 4927.0 25947.0 25545.0 27559.0

ITA 22950.0 10312.0 5718.0 4609.0 3100.0 24539.0

HUN 2175.0 4546.0 4142.0 4049.0 4091.0 2169.0 5260.0

ESP 6729.0 29126.0 27679.0 26004.0 24360.0 6714.0 4371.0 30033.0

POL 4616.0 22216.0 22335.0 21200.0 19573.0 4588.0 3933.0 22427.0 23828.0