Form GE Status as of: 2017-01-16

DESCRIPTION OF NATIONAL GENETIC EVALUATION SYSTEMS

SLOVENIJA Country (or countries) WORKABILITY Main trait group¹ **NOTE!** Only one trait group per form! HOL, BSW Breed(s) Milking Speed: linear score 1-5. Trait definition(s) and unit(s) of measurement² Attach an appendix if needed Only first lactation linear scoring is included in genetic evaluation. Method of measuring and collecting data Calving from 01.01.2004 Time period for data inclusion 1st parity Age groups (e.g. parities) included Classifier*year >= 10 scores Other criteria (data edits) for Classifier (total) \geq 20 scores inclusion of records (calving date - birth date) <= 1200 days 5 <= (scoring date - calving date) <= 365 Criteria for extension of records (if applicable) ΑI Sire categories No Environmental effects³, preadjustments ST - AM - BLUPMethod (model) of genetic evaluation³ F – classifier*year Environmental effects³ in the F – class(birth year)*class(scoring date - calving date) genetic evaluation model F – calving season*year R - herd*year Adjustment for heterogeneous variance in evaluation model Use of genetic groups and relationships No Blending of foreign/Interbull information in evaluation Use Appendix GE for heritability/genetic variance estimates; for multiple-trait genetic Genetic parameters in the evaluations, provide genetic correlation estimates between traits separately. evaluation Use also appendices PR, CO, BCO, SM, LO, CA, as applicable, if you participate in the international genetic evaluations of Interbull **System validation** Genetic trend validation – method 3 **Expression of genetic evaluations** BV12=((BV-a)/b)*12+100If standardised (e.g. RBV), give a - mean of BV standardisation formula in the b - standard deviation of BV appendix

Definition of genetic reference base	Population average					
Next base change	Next evaluation					
Calculation of reliability	Yes					
Criteria for official publication of evaluations	reliability >= 0.3					
Number of evaluations / publications per year	3					
Use in total merit index ⁴	HOL: Milking Speed 2%					
	BSW: Milking Speed 3%					
Anticipated changes in the near future						
Key reference on methodology applied	Web site: http://rodica.bf.uni-lj.si/govedo					
Key organisation: name, address, phone, fax, e-mail, web site	University of Ljubljana, Biotechnical Faculty, Department of Animal Science, Groblje 3, 1230 Domzale, Slovenija					
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¹⁾ Either: Production (e.g. milk, fat, protein), Conformation, Health (e.g. mastitis resistance, milk somatic cell, resistance to diseases other than mastitis), Longevity, Calving (e.g. stillbirth, calving ease), Female fertility (e.g. non-return rate, interval between reproductive events, number of AI's, heat strength), Workability (e.g. milking speed, temperament), Beef production, Efficiency (e.g. body weight, energy balance, body conditioning score), or Other traits.

²⁾ Indicate frequencies per category if the trait is categorical and specify transformation of data if practiced.

³⁾ Use abbreviations for most common effects (see document with list of abbreviations at http://www-interbull.slu.se/service_documentation/General/list_of_abbreviations.rtf) and indicate random (R) or fixed (F).

⁴⁾ Please give economic weights and indicate how they are expressed (preferably in genetic standard deviation units).

Form GE Appendix WO

Parameters for national genetic evaluations for workability traits as provided to Interbull

Country (or countries): SLOVENIJA

Main trait group: WORKABILITY

Breed (repeat as necessary): HOL

Trait	Definition	ITB ^a	h^{2b}	genetic variance ^b	official proof standardisation formula ^c
Milking Speed	Scale 1-5	X	0.088	0.040	a=0 b=0.0754 c=12 d=100
Temperament					

Country (or countries): SLOVENIJA

Main trait group: WORKABILITY

Breed (repeat as necessary): BSW

Trait	Definition	ITB ^a	h^{2b}	genetic variance ^b	official proof standardisation formula ^c
Milking Speed	Scale 1-5	X	0.103	0.046	a=0 b=0.0852 c=12 d=100
Temperament					

^a Indicate, with X, traits that are submitted to Interbull for international genetic evaluations.

b If repeated records are treated as separate traits, provide heritability estimates and genetic variances separately for each trait, as well as for all traits pooled, i.e. for the trait submitted to Interbull.

^c Expressed as follows: StandEval=((eval-a)/b)*c+d where a=mean of the base adjustment, b=standard deviation of the base, c=standard deviation of expression (include sign if scale is reversed), and d=base of expression.