Form GE Status as of: 2014-08-25

Updated April 24, 2007

DESCRIPTION OF NATIONAL GENETIC EVALUATION SYSTEMS

| Country (or countries) | SLOVENIJA |
|--|---|
| Main trait group ¹ NOTE! Only one trait group per | PRODUCTION |
| form! | HOL GDA DCW |
| Breed(s) | HOL, SIM, BSW |
| Trait definition(s) and unit(s) of measurement ² Attach an appendix if needed | Test day observations for milk (kg), protein (kg) and fat (kg) |
| Method of measuring and | ICAR Milk recording method to 29.2.2004 A4 method |
| collecting data | from 1.3.2004 AT4 method |
| Time period for data inclusion | Calving from 01.01.2004 |
| Age groups (e.g. parities) included | 1-5 parity |
| Other criteria (data edits) for | Days in milk 6-305. |
| inclusion of records | Number of herds >= 3. |
| Criteria for extension of records (if applicable) | |
| Sire categories | AI |
| Environmental effects ³ , pre- adjustments | No |
| Method (model) of genetic evaluation ³ | ST Repeatability TD |
| Environmental effects ³ in the genetic evaluation model | F – calving season, parity R – herd*year, permanent environment FR – Ali, Schaeffer |
| Adjustment for heterogeneous variance in evaluation model | |
| Use of genetic groups and relationships | |
| Blending of foreign/Interbull information in evaluation | No |
| Genetic parameters in the evaluation | Use Appendix GE for heritability/genetic variance estimates; for multiple-trait genetic evaluations, provide genetic correlation estimates between traits separately. |
| | 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+100 |
| If standardised (e.g. RBV), give standardisation formula in the | a – mean of BV |
| appendix | b – standard deviation of BV |
| Definition of genetic reference base | Mean of cows born in 2010 |
| Next base change | 2016 |
| Calculation of reliability | Yes |
| Interbull Code of Practice | Appendix II: Forms |
| Undeted April 24, 2007 | Appendix II. I offits |

| Criteria for official publication of evaluations | reliability(sire) >= 0.5, reliability(dam) >= 0.4 |
|---|---|
| Number of evaluations / publications per year | 3 |
| Use in total merit index ⁴ | HOL: Fat yield 4%, protein yield 16% |
| | BSW: Fat yield 4.5%, protein yield 17% |
| | SIM: Fat yield 5%, protein yield 20% |
| Anticipated changes in the near future | Change of genetic base |
| Key reference on methodology applied | Web site: http://www.bf.uni-lj.si/zootehnika/struktura/katedre-in-enote/center-za-strokovno-delo-v-zivinoreji/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 Tel. +386 1 3203 872 Fax: +386 1 7241 005 Jurij.Krsnik@bf.uni-lj.si, Klemen.Potocnik@bf.uni-lj.si |

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://wwwinterbull.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 GE

Parameters used in genetic evaluation

Country (or countries): SLOVENIJA

Main trait group: PRODUCTION

Breed (repeat as necessary): HOL

| Trait | Definition | ITB ^a | h ^{2b} | genetic variance ^b | official proof standardisation formula ^c |
|---------|-----------------|------------------|-----------------|----------------------------------|--|
| Milk | TD record in kg | X | 0.25 | 10.708 kg^2 | BV12=((BV-(0.9291))/2.1399)*12+100 |
| Fat | TD record in kg | X | 0.19 | $0.016\mathrm{kg^2}$ | BV12=((BV-(0.0359))/0.0819)*12+100 |
| Protein | TD record in kg | X | 0.21 | $0.01~\mathrm{kg}^2$ | BV12=((BV-(0.0348))/0.0627)*12+100 |

Country (or countries): SLOVENIJA

Main trait group: PRODUCTION

Breed (repeat as necessary): SIM

| Trait | Definition | ITB ^a | h ^{2b} | genetic variance ^b | official proof standardisation formula ^c |
|---------|-----------------|------------------|-----------------|----------------------------------|--|
| Milk | TD record in kg | X | 0.38 | 9.309 kg^2 | BV12=((BV-(1.1229))/2.4415)*12+100 |
| Fat | TD record in kg | X | 0.299 | $0.016~\mathrm{kg^2}$ | BV12=((BV-(0.0448))/0.0985)*12+100 |
| Protein | TD record in kg | X | 0.308 | $0.009~\mathrm{kg^2}$ | BV12=((BV-(0.0325))/0.0712)*12+100 |

Country (or countries): SLOVENIJA

Main trait group: PRODUCTION

Breed (repeat as necessary): BSW

| Trait | Definition | ITB ^a | h ^{2b} | genetic variance ^b | official proof standardisation formula ^c |
|---------|-----------------|------------------|-----------------|----------------------------------|---|
| Milk | TD record in kg | X | 0.32 | $7.756~\mathrm{kg}^2$ | BV12=((BV-(1.0938))/2.0097)*12+100 |
| Fat | TD record in kg | X | 0.23 | $0.012~\mathrm{kg}^2$ | BV12=((BV-(-0.0436))/0.0788)*12+100 |
| Protein | TD record in kg | X | 0.25 | $0.007~\mathrm{kg}^2$ | BV12=((BV-(0.0401))/0.0618)*12+100 |

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

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.

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.

Form GE Appendix PR

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

Country (or countries): SLOVENIJA

Main trait group: Production

Breed(s): HOL

| Trait | h^{2a} | Genetic variance ^a | official proof standardisation formula ^b |
|----------------|----------|-------------------------------|--|
| Milk yield: | 0.25 | $10.708~\mathrm{kg}^2$ | BV12=((BV-(0.9291))/2.1399)*12+100 |
| Fat yield: | 0.19 | $0.016\mathrm{kg}^2$ | BV12=((BV-(0.0359))/0.0819)*12+100 |
| Protein yield: | 0.21 | $0.01~\mathrm{kg}^2$ | BV12=((BV-(0.0348))/0.0627)*12+100 |

Country (or countries): SLOVENIJA

Main trait group: Production

Breed(s): SIM

| Trait | h^{2a} | Genetic variance ^a | official proof standardisation formula ^b |
|----------------|----------|-------------------------------|--|
| Milk yield: | 0.38 | $9.309~\mathrm{kg}^2$ | BV12=((BV-(1.1229))/2.4415)*12+100 |
| Fat yield: | 0.299 | $0.016\mathrm{kg}^2$ | BV12=((BV-(0.0448))/0.0985)*12+100 |
| Protein yield: | 0.308 | $0.009~\mathrm{kg}^2$ | BV12=((BV-(0.0325))/0.0712)*12+100 |

Country (or countries): SLOVENIJA

Main trait group: Production

Breed(s): BSW

| Trait | $ m h^{2a}$ | Genetic variance ^a | official proof standardisation formula ^b |
|----------------|-------------|-------------------------------|--|
| Milk yield: | 0.32 | $7.756\mathrm{kg}^2$ | BV12=((BV-(1.0938))/2.0097)*12+100 |
| Fat yield: | 0.23 | $0.012~\mathrm{kg}^2$ | BV12=((BV-(-0.0436))/0.0788)*12+100 |
| Protein yield: | 0.25 | $0.007~\mathrm{kg}^2$ | BV12=((BV-(0.0401))/0.0618)*12+100 |

- ^a If lactations, or part of lactations, are treated as separate traits, provide heritability estimates and genetic variances separately for each lactation, as well as for all lactations pooled, i.e. for the trait submitted to Interbull.
- 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.