

**Country**

**Germany**

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**Trait category:**

**Individual trait(s):**

**Reproduction-calving**

**Calving performance (direct, maternal)**

**Stillbirth (direct, maternal)**

**Reproduction-fertility**

**Non-return rate 90 (female, male)**

**Health**

**Somatic cell count**

**Workability**

**Milking speed**

**Conformation**

**Udder**

**Locomotion**

**Other**

**Longevity**

**Stayability**

**Coordination:**

**Arbeitsgemeinschaft Deutscher Rinderzüchter E.V. (A.D.R.)**

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## GERMANY

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Addresses of the different co-operatives responsible for evaluation and publication for different breeds/traits:

All evaluation for Holstein Friesian and German Red & White, except for stayability, milking speed and beef & growth traits:

Vereinigte Informationssysteme Tierhaltung w.V. (VIT Verden)

Heideweg 1

D-27283 Verden, Germany

Telephone +49 4231 955 - 171 or 173

Facsimile +49 4231 955 - 166

E-mail vitall@rzv-srv.vit.de

All breeds restricted to Schleswig-Holstein for somatic cell count and stayability:

LKV Schleswig-Holstein

Steenbeker Weg 151

D-24106 Kiel, Germany

Telephone +49 431 339870

Facsimile +49 431 3398713

All breeds restricted to Rheinland-Pfalz and Saarland for milking speed:

Landesanstalt für Tierzucht und Qualitätsprüfungen Neumühle

D-67728 Münchweiler/Alsenz, Germany

Telephone +49 6302 92160

Facsimile +49 6302 921699

All breeds restricted to Bavaria for somatic cell count and beef & growth traits:

Landeskuratorium der Erzeugerringe für Tierische Veredlung

Haydnstr. 11

D-80336 München, Germany

Telephone +49 89 5443480

Facsimile +49 89 54434810

All breeds restricted to Bavaria, except for somatic cell count and beef & growth traits:

Bayerische Landesanstalt für Tierzucht

Prof. Dürrwaechter-Platz 1

D-85586 Poing, Germany

Telephone +49 89 99141 300

Facsimile +49 89 99141 105

All breeds restricted to Baden-Württemberg:

Landesamt für Flurneuordnung und Landesentwicklung

EB21

D-70806 Kornwestheim

Telephone +49 7154 139368

Facsimile +49 7154 139499

| Reproduction<br>calving traits                              | Calving performance (direct, maternal)<br>Stillbirth (direct, maternal) |  |
|---|---|--|
| Breed(s)  | [V]<br>[B]  | Verden: Holstein Friesian, German Red & White<br>Bavaria: Fleckvieh, Braunvieh, Gelbvieh   |
| Trait definition and<br>unit(s) of measuring                | [V]<br><br>[B]  | Calving performance is scored in 3 categories;<br>normal or easy (1), hard pull (2), veterinary help or<br>caesarian (3)<br>Stillbirth is defined as stillborn or died within 24<br>hours after birth<br>Calving performance is scored in 5 categories;<br>without help (0), one helper (1), two or more<br>helpers or mechanical help (3), veterinary help (4),<br>caesarian section (5)<br>Stillbirth is defined as stillborn or died within 48<br>hours after birth |
| Method of measuring and<br>collecting data                  | [V,B]   | Scored by farmer and collected by milk recording<br>system   |
| Time period for data<br>inclusion                           | [V]<br>[B]  | Since 1986<br>Since 1979   |
| Age groups  | [V,B]   | All  |
| Genetic parameters  | [V,B]   | $h^2_{\text{calving performance (direct)}} = 0.05$<br>$h^2_{\text{calving performance (maternal)}} = 0.05$<br>$r_g(\text{calving performance (direct, maternal)}) = -0.10$<br>$h^2_{\text{stillbirth (direct)}} = 0.05$<br>$h^2_{\text{stillbirth (maternal)}} = 0.05$<br>$r_g(\text{stillbirth (direct, maternal)}) = -0.10$  |
| Sire categories   | [V,B]   | All bulls  |
| Environmental effects<br>pre-adjustment<br>evaluation model | [V,B]<br>[V,B]  | None<br>Herd x calving year, calving month, parity, age<br>within parity, sex of calf, permanent cow effect  |
| Base for age adjustment                                     | [V,B]   | None   |
| Use of genetic groups<br>and/or relationships               | [V,B]   | AM with genetic "phantom" groups for unknown<br>parents  |
| Method (model) of genetic<br>evaluation                     | [V,B]   | Maternal effects ST BLUP AM for repeated<br>records  |
| System validation   | [V,B]   | Plausibility checks of recorded data, check on<br>model suitability (fixed effects, EBV), genetic<br>trend   |
| Expression of proof   | [V,B]   | RBV-C standardized with M = 100 and SD = 12,<br>higher values are more desirable   |
| Genetic (reference) base                                    | [V,B]   | All cows born 1990 within breed  |
| Criteria for official<br>publication of sire proofs         | [V]<br>[B]  | REL > 0.50<br>≥ 10 daughters in ≥ 5 farms for milk production  |

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| Reproduction<br>calving traits <i>continued</i> | Calving performance (direct, maternal)<br>Stillbirth (direct, maternal)   |
|---|---|
| Number of evaluations/<br>publications per year | [V] One; August<br>[B] Two; May, November   |
| Use in total merit index                        | [V,B] Development of selection index in progress  |
| Key reference on<br>methodology applied         | [V,B] Gierdziewicz, M. et al., 1994. 45 <sup>th</sup> Meeting of<br>EAAP, Edinburgh. Evaluation of calving ease using<br>a reduced animal model in German Fleckvieh<br>Averdunk, G. et al., 1995. Proc. of the open<br>session of the Interbull annual meeting Prague. Sire<br>evaluation for fertility and calving ease in Germany |

| Reproduction fertility traits                         | Non-return rate 90 (female, male) |   |
|---|-----------------------------------|---|
| Breed(s)  | [V]<br>[B]<br>[W]                 | Verden: Holstein Friesian, German Red & White<br>Bavaria: Fleckvieh, Braunvieh, Gelbvieh<br>Baden-Württemberg: Fleckvieh, Braunvieh, Vorderwälder, Schwarzbunt, Rotbunt |
| Trait definition and unit(s) of measuring             | [V,B,W]                           | Percentage non-returns within 90 days after first insemination  |
| Method of measuring and collecting data               | [V]                               | Collected by AI-service technicians, veterinarians and milk recording system  |
|   | [B,W]                             | Collected by AI-organizations   |
| Time period for data inclusion                        | [V,B]                             | Since 1986  |
|   | [W]                               | Since 1980  |
| Age groups  | [V,B,W]                           | All   |
| Genetic parameters                                    | [V,B,W]                           | $h^2_{\text{non-return rate 90 (female)}} = 0.02$<br>$h^2_{\text{non-return rate 90 (male)}} = 0.02$<br>$r_{g(\text{non-return rate 90 (female, male)})} = -0.05$       |
| Sire categories                                       | [V,B,W]                           | All bulls   |
| Environmental effects pre-adjustment evaluation model | [V,B,W]                           | None  |
|   | [V,B,W]                           | Herd x year, calving month, parity, age within parity (only heifers), interval between calving and 1 <sup>st</sup> insemination (only cows), permanent cow effect       |
| Base for age adjustment                               | [V,B,W]                           | None  |
| Use of genetic groups and/or relationships            | [V,B,W]                           | AM with genetic "phantom" groups for unknown parents  |
| Method (model) of genetic evaluation                  | [V,B,W]                           | ST BLUP AM, including paternal and maternal genetic effects   |
| System validation                                     | [V,B,W]                           | Plausibility checks of recorded data, checks on model suitability for fixed effects and breeding value  |
| Expression of proof                                   | [V,B,W]                           | RBV-F standardized with $M = 100$ and $SD = 12$ , higher values are more desirable  |
| Genetic (reference) base                              | [V,B]                             | All cows born in 1990 within breed  |
|   | [W]                               | Fixed, sires born in 1982-1984  |
| Criteria for official publication of sire proofs      | [V]                               | REL > 0.50  |
|   | [B]                               | ≥ 10 daughters in ≥ 5 herds for milk production   |
|   | [W]                               | ≥ 200 insemination and ≥ 50 daughters inseminations   |
| Number of evaluations/publications per year           | [V]                               | One; August   |
|   | [B]                               | Two; May, November  |
| Use in total merit index                              | [V,B]                             | Development of selection index in progress  |

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### Reproduction

fertility traits *continued*

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### Non-return rate 90 (female, male)

#### Key reference on methodology applied

- [V,B] Thaller, G et al., 1994. 45<sup>th</sup> Meeting of EAAP, Edinburgh. Breeding value estimation for reproductive traits by an animal model with paternal and maternal effects
- Averdunk, G. et al., 1995. Proc. of the open session of the Interbull annual meeting Prague. Sire evaluation for fertility and calving ease in Germany

| Health traits   | Somatic cell count |  |
|---|--------------------|--|
| Breed(s)  | [V]                | Verden: Holstein Friesian, German Red & White  |
|   | [B]                | Bavaria: Fleckvieh, Braunvieh, Gelbvieh, Holstein Friesian   |
|   | [S]                | Schleswig Holstein: Holstein Friesian, German Red & White, Red Angler  |
| Trait definition and unit(s) of measuring             | [V]                | Somatic cell score is log 2 transformed test-day somatic cell counts, taken between 4 and 365 days in lactation; number of cells should be between 5,000 and 6,400,000 |
|   | [B]                | Log 10 transformed somatic cell counts (in 1000/ml); number of cells should be between 10,000 and 10,000,000   |
|   | [S]                | Lactation mean of log transformed test-day somatic cell count, taken between 2 <sup>nd</sup> and 8 <sup>th</sup> milk recording  |
| Method of measuring and collecting data               | [V,B]              | Test-day results from milkrecording  |
|   | [S]                | Collected during normal milk recording   |
| Time period for data inclusion                        | [V]                | Since 1990   |
|   | [B]                | Since 1989   |
|   | [S]                | Since 1986   |
| Age groups  | [V]                | 1 <sup>st</sup> to 3 <sup>rd</sup> lactation   |
|   | [B]                | 1 <sup>st</sup> lactation  |
|   | [S]                | All  |
| Genetic parameters                                    | [V]                | $h^2_{\text{somatic cell score (lactation 1)}} = 0.08$   |
|   |                    | $h^2_{\text{somatic cell score (lactation 2)}} = 0.13$   |
|   |                    | $h^2_{\text{somatic cell score (lactation 3)}} = 0.14$   |
|   |                    | $r_g(\text{somatic cell score (lactation 1, lactation 2)}) = 0.90$   |
|   |                    | $r_g(\text{somatic cell score (lactation 1, lactation 3)}) = 0.85$   |
|   | [B]                | $r_g(\text{somatic cell score (lactation 2, lactation 3)}) = 0.97$   |
|   |                    | $h^2_{\text{somatic cell count (for Fleckvieh and Gelbvieh)}} = 0.05$  |
|   |                    | $h^2_{\text{somatic cell count (for Braunvieh and Holstein Friesian)}} = 0.07$   |
|   | [S]                | $h^2_{\text{somatic cell count}} = 0.10$   |
| Sire categories                                       | [V,S]              | All bulls  |
|   | [B]                | Bulls with $\geq 10$ daughters   |
| Environmental effects pre-adjustment evaluation model | [V,B,S]            | None   |
|   | [V]                | Region, age of calving, stage of lactation, season of calving, herd x test-date, permanent environment   |
|   | [B]                | Herd x year, month, stage of lactation, cow  |
|   | [S]                | Herd x year, parity  |
| Base for age adjustment                               | [V,B,S]            | None   |
| Use of genetic groups and/or relationships            | [V]                | All known relations in an AM. Phantom parent groups according to selection path, breed, year of birth  |
|   | [B]                | None   |
|   | [S]                | Bull's birth year  |

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| Health traits <i>continued</i>                   |         | Somatic cell count  |
|--|---------|---|
| Method (model) of genetic evaluation             | [V]     | MT BLUP AM for test-day records. Lactation 1, 2 and 3 are considered as separate traits   |
|  | [B,S]   | ST BLUP Sire repeatability model  |
| System validation                                | [V]     | Edits for age of calving and days in milk   |
|  | [B]     | Checks similar to milk recorded data  |
|  | [S]     | Plausibility checks of recorded data  |
| Expression of proof                              | [V]     | Combined RBV = $0.26 \times \text{scs}_{\text{lactation 1}} + 0.37 \times \text{scs}_{\text{lactation 2}} + 0.37 \times \text{scs}_{\text{lactation 3}}$ , with M = 100 and SD = 12, higher values are more desirable   |
|  | [B,S]   | RBV standardized with M = 100 and SD = 12, higher values are more desirable   |
| Genetic (reference) base                         | [V]     | Rolling bull base, defined by all 8-10 year old AI-bulls  |
|  | [B]     | All test bulls tested in the latest 3 year period for milk production   |
|  | [S]     | Rolling bull base   |
| Criteria for official publication of sire proofs | [V,B,S] | REL $\geq$ 50 %   |
| Number of evaluations/publications per year      | [V]     | Two; March, September   |
|  | [B]     | Two; April, October   |
|  | [S]     | One; August   |
| Use in total merit index                         | [V,B,S] | No  |
| Key reference on methodology applied             | [V]     | Reents, R., J. Jamrozik, L.R. Schaeffer & J.C.M. Dekkers, 1995. Estimation of genetic parameters for test-day records of somatic cell score. J. Dairy Sci. 78: 2847<br>Reents, R., J.C.M. Dekkers & L.R. Schaeffer, 1995. Genetic evaluation for somatic cell score with a test-day model for multiple lactations. J. Dairy Sci. 78: 2858 |



| Workability traits                               | Milking speed     |  |
|--|-------------------|--|
| Breed(s)   | [B]<br>[R]<br>[W] | Bavaria: Fleckvieh, Braunvieh, Gelbvieh<br>Rheinland-Pfalz and Saarland: Holstein Friesian, German Red & White, Red Angler<br>Baden-Württemberg: Fleckvieh, Braunvieh, Vorderwälder, Schwarz & Rotbunt |
| Trait definition and unit(s) of measuring        | [B,W]<br>[R]      | Average milking speed (in kg/minute)<br>Average milking speed measured between 50 <sup>th</sup> and 180 <sup>th</sup> day during first lactation (in kg/minute)  |
| Method of measuring and collecting data          | [B]<br>[R,W]      | Included in milk recording scheme<br>Collected during normal milk recording  |
| Time period for data inclusion                   | [B]<br>[R]<br>[W] | Since 1970<br>Since 1976<br>Since 1981   |
| Age groups                                       | [B,R,W]           | 1 <sup>st</sup> lactation  |
| Genetic parameters                               | [B,R]<br>[W]      | $h^2_{\text{milking speed}} = 0.20$<br>$h^2_{\text{milking speed}} = 0.30$   |
| Sire categories                                  | [B,W]<br>[R]      | Test bulls<br>All bulls with progeny records   |
| Environmental effects pre-adjustment             | [B]<br>[R]<br>[W] | Test-day milk yield<br>Test-day within lactation, herd, year<br>Day of lactation   |
| evaluation model                                 | [B]<br>[R]        | Technician doing the proof<br>None   |
| Base for age adjustment                          | [B,R,W]           | None   |
| Use of genetic groups and/or relationships       | [B,R]<br>[W]      | None<br>Birth year of sire   |
| Method (model) of genetic evaluation             | [B,W]<br>[R]      | ST CC-procedure<br>ST SM   |
| System validation                                | [B]<br>[R]<br>[W] | Plausible checks of recorded data<br>Check for extreme values (excess, kurtosis)<br>-  |
| Expression of proof                              | [B,R,W]           | EBV in kg milk per minute  |
| Genetic (reference) base                         | [R]<br>[W]        | All bulls proved within the last 5 years<br>Bulls born 1982-1984   |
| Criteria for official publication of sire proofs | [B,R]<br>[W]      | $\geq 20$ daughters<br>$\geq 20$ daughters (for Fleckvieh)<br>$\geq 10$ daughters (for other breeds)   |
| Number of evaluations/publications per year      | [B]<br>[R]        | Four; February, May, August, November<br>Two; June, December   |
| Use in total merit index                         | [B]<br>[R,W]      | Development of selection index in progress<br>No   |
| Key reference on methodology applied             | [B,R,W]           | -  |

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|   |                    |  |
|---|--------------------|--|
| <b>Conformation traits</b>                        | <b>Udder:</b>      | fore udder attachment, fore udder length, rear udder height, suspensory ligament, udder depth, teat placement, teat length, udder overall  |
|   | <b>Locomotion:</b> | rear leg set, foot angle   |
|   | <b>Other:</b>      | stature, body depth, rump angle, rump width, strength, dairy character, body type overall  |
| <b>Breed(s)</b>                                   | [V]                | Verden: Holstein Friesian, German Red & White, Angler  |
|   | [B]                | Bavaria: Fleckvieh, Braunvieh, Gelbvieh  |
|   | [W]                | Baden-Württemberg: Fleckvieh, Braunvieh, Vorderwälder, Schwarz & Rotbunt   |
| <b>Trait definition and unit(s) of measuring</b>  | [V]                | Most individual traits scored on a linear 1-9 point scale, following recommendation of the European and World-wide group for harmonization of linear type classification, except for stature, which is measured in cm<br>Overall traits are scored on a 0-50 point scale |
| <b>Method of measuring and collecting data</b>    | [V]                | Scored by classifier   |
|   | [B,W]              | Scored by official people  |
| <b>Time period for data inclusion</b>             | [V]                | Since 1984   |
|   | [B,W]              | New scoring system since 1988  |
| <b>Age groups</b>                                 | [V,B,W]            | 1 <sup>st</sup> lactation  |
| <b>Genetic parameters</b>                         | [V]                | $h^2_{\text{udder traits}} = 0.18 \text{ to } 0.27$<br>$h^2_{\text{locomotion traits}} = 0.13$<br>$h^2_{\text{other traits}} = 0.21 \text{ to } 0.43$  |
| <b>Sire categories</b>                            | [V]                | All bulls  |
|   | [B]                | Promising AI-sires, since 1993 all AI-sires  |
|   | [W]                | Promising AI-sires   |
| <b>Environmental effects pre-adjustment</b>       | [V]                | Heterogeneous variances between classifiers  |
|   | [B,W]              | None   |
| <b>evaluation model</b>                           | [V]                | Classifier x year, herd x year, stage of lactation, age of calving   |
|   | [B]                | Classifier   |
|   | [W]                | Time x region x person   |
| <b>Base for age adjustment</b>                    | [V,B,W]            | None   |
| <b>Use of genetic groups and/or relationships</b> | [V]                | All known relations in an Animal model. Phantom parent groups according to selection path, breed, year of birth  |
|   | [B,W]              | None   |
| <b>Method (model) of genetic evaluation</b>       | [V]                | ST BLUP AM   |
|   | [B]                | ST deviation of average of classifier  |
|   | [W]                | CC-procedure   |

| Conformation traits<br><i>continued</i>          | Udder      |  |
|--|------------|--|
|  | Locomotion | Other  |
| System validation                                | [V]        | Check for reasonable age of calving and stage of lactation at time of classification, only 1 <sup>st</sup> lactation 1 <sup>st</sup> classification are used |
|  | [B,W]      | None   |
| Expression of proof                              | [V]        | EBV standardized with M = 100 and SD = 12  |
|  | [B]        | RBV with M = 100 and SD = 12, higher values are more desirable   |
|  | [W]        | EBV  |
| Genetic (reference) base                         | [V]        | Rolling bull base, defined by all 8-10 year old AI-bulls   |
|  | [B]        | None   |
| Criteria for official publication of sire proofs | [V]        | ≥ 20 daughters in 5 herds  |
|  | [B]        | ≥ 30 daughters   |
|  | [W]        | ≥ 20 daughters   |
| Number of evaluations/publications per year      | [V]        | Two; March, September  |
|  | [B]        | Four; February, May, August, November  |
| Use in total merit index                         | [V,R,B]    | No   |
| Key reference on methodology applied             | [V,B]      | Reents, R, 1993. Estimation of breeding values for type traits in Germany, Interbull Bulletin No. 8, 1993  |

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| Longevity traits                                      | Stayability           |  |
|---|-----------------------|--|
| Breed(s)  | [B]<br>[S]<br>[W]     | Bavaria: Fleckvieh, Braunvieh, Gelbvieh<br>Schleswig Holstein: Holstein Friesian, German Red & White, Red Angler<br>Baden-Württemberg: Fleckvieh, Braunvieh, Vorderwälder, Schwarz & Rotbunt |
| Trait definition and unit(s) of measuring             | [B]<br>[S,W]          | Relative risk for culling<br>Percentage of live daughters at the age of 46 and 60 months   |
| Method of measuring and collecting data               | [B]<br>[S]<br>[W]     | Included in milk recording scheme<br>Birth and culling date of cows recorded during normal milk recording<br>Birth dates of all cows   |
| Time period for data inclusion                        | [B]<br>[S]<br>[W]     | Since 1976<br>Since 1979<br>Since 1971   |
| Age groups  | [B,S,W]               | All  |
| Genetic parameters                                    | [B]<br>[S]<br>[W]     | $h^2_{\text{stayability}} = 0.1$<br>$h^2_{\text{stayability}} = 0.05$<br>$h^2_{\text{stayability}} = 0.07 \text{ to } 0.12$<br>(different $h^2$ for different breeds)                        |
| Sire categories                                       | [B,S,W]               | All bulls  |
| Environmental effects pre-adjustment evaluation model | [B,S,W]<br>[S]<br>[W] | None<br>Herd, year x season, age of first calving<br>Herd, first calving age group, genetic group  |
| Base for age adjustment                               | [B,S,W]               | None   |
| Use of genetic groups and/or relationships            | [B]<br>[S]<br>[W]     | Full relationship matrix, animals with unknown parents are grouped by birth year<br>Bull's birth year<br>Birth year of sire of semen   |
| Method (model) of genetic evaluation                  | [B,S,W]               | ST BLUP SM   |
| System validation                                     | [B,W]<br>[S]          | -<br>Plausibility checks of recorded data  |
| Expression of proof                                   | [S]<br>[W]            | RBV standardized with $M = 100$ and $SD = 12$ , higher values are more desirable<br>EBV  |
| Genetic (reference) base                              | [B]<br>[S]<br>[W]     | Rolling bull base, i.e. in 1996 average of 1986 to 1988 was used<br>Rolling bull base, i.e. in 1996 average of 1983 to 1986 was used<br>Bulls born between 1975-1977                         |
| Criteria for official publication of sire proofs      | [S]<br>[W]            | $REL > 50\%$<br>$\geq 50$ daughters (for Fleckvieh)<br>$\geq 30$ daughters (for other breeds)  |

**Longevity traits****Stayability***continued***Number of evaluations/  
publications per year**

[B]

Two; May, November

[S]

One; August

**Use in total merit index**

[B,S,W]

No

**Key reference on  
methodology applied**

[B,S,W]

-