



Genetic Analysis of Reproductive Disorders and their Relationship to Fertility in Fleckvieh Dual Purpose Cattle



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- **Decreased fertility main reason for involuntary culling of dairy cows**
- **Fertility is a biologically complex trait**
 - ” Recycle after calving
 - ” Show heat
 - ” Successful insemination
 - ” Embryo survival
 - ” Resistance to reproductive disorders
- **Fertility index**
 - ” Non-return rate 56 (NR56)
 - ” Interval from calving to first insemination (CFI)
 - ” Interval from first to last insemination (FLI)

- **Denmark, Sweden and Finland consider reproductive disorders**
- **Since July 2006 Health monitoring in cattle in Austria**

Recording of diagnoses data from veterinarians

Data used for herd management and development of genetic evaluation for health traits



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Objectives



- **Estimate heritabilities for various reproductive disorders with linear and threshold models**
- **Investigate relationships between reproductive disorders and fertility traits (NR56, CFI and FLI)**

- **Austrian Fleckvieh dual purpose cattle**
- **Herds with reliable health recording**
- **Calvings between Jan. 2007 and Nov. 2008**
- **All cows from parity 1 to 5**

- **Metritis (MET)**
 - **Silent heat and anoestrus (ESTRUS)**
 - **Cystic ovaries (CYST)**
 - **Retained placenta (RP)**
 - **Puerperal diseases (PUERP)**
 - **Uterine prolapse**
 - **Abortion**
 - **Difficult calving**
 - **Calving injury**
- } Disease frequencies far
below 1 %

Trait	Days after calving	Records	Mean
MET¹	0 to 150 d	13,958	3.8%
ESTRUS¹	0 to 150 d	13,958	6.3%
CYST¹	0 to 150 d	13,958	9.6%
RP	0 to 7 d	22,355	2.6%
PUERP	0 to 30 d	20,986	2.5%
MET/RP/PUERP	0 to 30 d	20,986	6.4%
REPRO²	0 to 150 d	15,620	15.7%
NR56		20,578	66.4%
CFI		24,725	69.6 d
FLI		11,262	23.4 d

¹ Only cows that completed at least two-thirds of the opportunity period (100 d)

² MET + CYST + RP + PUERP + culling reason included

➤ Heritabilities

- ” Linear animal model, VCE
- ” Threshold sire model (logit link), ASReml

➤ Genetic correlations

Linear animal model, bivariate analyses, VCE

reproductive disorders

$$y_{ijklm} = \mu + L_i + HYS_j + a_k + pe_l + e_{ijklm}$$

- y_{ijklm} = individual observation (0/1)
- μ = overall mean
- L_i = fixed effect of parity
- HYS_j = random effect of herd*year*season of calving
- a_k = random additive genetic effect of animal or sire
- pe_l = random permanent environmental effect of cow
- e_{ijklm} = residual error term

ility traits

$$y_{ijklmn} = \mu + YM_i + L_j + HY_k + a_l + pe_m + e_{ijklmn}$$

- y_{ijklmn} = individual observation
- μ = overall mean
- YM_i = fixed effect of year*month of calving or insemination
- L_j = fixed effect of calving age-parity
- HY_k = random effect of herd*year of calving or insemination
- a_l = random additive genetic effect of animal
- pe_m = random permanent environmental effect of cow
- e_{ijklmn} = residual error term

reproductibilities

Trait	Linear animal model	Threshold sire model
MET	0.01 _{0.004}	0.06 _{0.036}
ESTRUS	0.01 _{0.004}	0.01 _{0.015}
CYST	0.04 _{0.009}	0.08 _{0.027}
RP	0.01 _{0.004}	0.06 _{0.038}
PUERP	0.02 _{0.005}	0.14 _{0.054}
MET/RP/PUERP	0.02 _{0.005}	0.08 _{0.027}
REPRO	0.04 _{0.009}	0.07 _{0.020}

Genetic correlations

Trait	MET	CYST	RP	PUERP
MET	0.01 (0.004)	0.22 (0.235)	1.00 (<0.001)	1.00 (0.001)
CYST		0.04 (0.009)	-0.01 (0.180)	0.68 (0.114)
RP			0.01 (0.004)	1.00 (<0.001)
PUERP				0.02 (0.005)

Genetic correlations

Trait	MET	CYST	RP	PUERP
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RP			0.01 (0.004)	1.00 (<0.001)
PUERP				0.02 (0.005)



Genetic correlations

Trait	MET/RP/ PUERP	CYST	REPRO	NR56	CFI	FLI
MET/RP/ PUERP	0.02 (0.005)	0.48 (0.137)	0.77 (0.085)	-0.37 (0.228)	0.26 (0.134)	0.77 (0.184)
CYST		0.04 (0.009)	0.90 (0.037)	-0.70 (0.141)	0.29 (0.166)	0.74 (0.137)
REPRO			0.04 (0.009)	-0.76 (0.211)	0.42 (0.157)	0.93 (0.071)
NR56				0.01 (0.004)	-0.23 (0.231)	-0.93 (0.132)
CFI					0.03 (0.007)	0.52 (0.176)
FLI						0.01 (0.006)



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- **Data from Austrian project BHealth monitoring in cattle can be used for genetic selection**
- **Sufficient genetic variation exists for all reproductive disorders, except ESTRUS**
- **Genetic correlations between reproductive disorders and fertility were favourable**
- **For routine genetic evaluation trait REPRO is recommended**



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