

Genetic correlations between female fertility and production traits in South African Holstein cattle

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Abstract

Female fertility is increasingly gaining importance in national dairy cattle breeding objectives worldwide. In South Africa, there is no routine prediction of breeding values for reproductive performance in dairy cattle and selection is mainly focused on production traits. The objective of this study was to estimate genetic parameters among female fertility traits (age at first calving and calving interval) and first, second and third lactation production traits in South African Holstein cattle to determine the effect that selection on production *per se* may have on female fertility. Performance records on 40 437 South African Holstein cows in 766 herds were used. (Co)variance estimates were obtained by multitrait analysis, using the REML procedure. Heritability estimates were moderate for age at first calving (0.24 ± 0.02) and low for calving interval (0.03 ± 0.01). Genetic correlations between age at first calving and yield traits were low to moderately negative, ranging from -0.17 ± 0.07 with second lactation butterfat percentage to -0.50 ± 0.05 with first lactation butterfat yield. Calving interval had moderate to highly positive genetic correlations with yield traits, ranging from 0.37 ± 0.10 with second lactation milk yield to 0.69 ± 0.06 with first lactation milk yield. Correlations between female fertility and butterfat and protein percentages across all lactations, were close to zero. The observed antagonistic relationship between calving interval and production traits highlights the need to include calving interval in breeding objectives for South African Holstein cattle.

Keywords: Age at first calving, calving interval, milk yield

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Introduction

Female fertility is one of the most economically important traits in dairy cattle. Reproductive failure causes economic losses due to reduced production as a result of prolonged calving intervals (Van Arendonk *et al.*, 1989; Boichard *et al.*, 1997; Olori *et al.*, 2002), increased insemination costs, reduced returns from calves born and higher replacements costs (Bagnato & Oltenacu, 1994). Fertility problems are the most common reasons for culling in dairy cattle (Philipsson, 1981). Culling for low reproduction rate as a proportion of disposal reasons has been reported to account for 25% in France (Colleau & Moureaux, 1999). Esselmont & Kossaibati (1997) showed that the primary reason for culling dairy cattle in the UK was failure to conceive, which accounted for 44% of culls in first lactation animals. Depending on the level of production, increasing calving interval (CI) by one day costs the breeder about 1.8 US dollars in Ireland, without accounting for the costs of higher culling due to poorer fertility (Esselmont *et al.*, 2001).

Female fertility has been neglected in the past in most dairy cattle genetic improvement programmes worldwide, mainly because fertility traits are known to exhibit a low heritability (Raheja *et al.*, 1989; Grosshans *et al.*, 1997; Pryce *et al.*, 1998; Kadarmideen, 2004). Despite their low heritability, fertility traits have been shown to have high additive genetic variation (Philipsson, 1981; Hermas *et al.*, 1987; Raheja *et al.*, 1989; Oltenacu, 1991; Grosshans *et al.*, 1997; De Jong, 1998) and therefore, increasing the amount of information available for use in genetic evaluation may facilitate their improvement through selection. Traditionally selection pressure has been applied mainly on yield traits worldwide. Because of antagonistic genetic correlations, intense single-trait selection for production is expected to cause a decline in fertility (Van Arendonk *et al.*, 1989; Frick & Lindhe, 1991; Bagnato & Oltenacu, 1994; Campos *et al.*, 1994; Hoekstra *et al.*, 1994; Pryce *et al.*, 1997; 2004; De Jong, 1998; Ojango & Pollot, 2001; Nilfroooshan & Edriss, 2004; Kadarmideen, 2004; VanRaden *et al.*, 2004).

To maintain or recover a high fertility in modern dairy cows calls for a two-pronged approach, involving both inclusion of fertility in broader breeding objectives and adjustment to management practices (Pryce *et al.*, 2004). Selection for fertility traits requires identifying suitable traits that can be used as selection criteria (Haile-Mariam *et al.*, 2003). Kadarmideen & Simm (2002) observed an increase in economic returns of up to 38% by adding CI to the UK total merit index.

Breeding programmes for dairy cattle in South Africa have been based primarily on increased milk production (Banga & Rautenbach, 1999). During the period 1982 to 2001, the average genetic merit for production traits increased remarkably in the major dairy cattle breeds (Hallowell & Mostert, 2001). These trends are probably due to the fact that yield is the main selection criterion for most South African producers.

The objectives of this study were: a) to estimate genetic parameters of female fertility traits (age at first calving (AFC) and CI) and, first, second and third lactation production traits; and b) to determine the effect that selection on production *per se* may have on female fertility.

Materials and Methods

The fertility traits considered were AFC and CI. Calving interval in different lactations was treated as the same repeated trait. Production traits were lactation milk yield (MLKY), butterfat yield (BFATY), protein yield (PROTY), butterfat percentage (BFPCT) and protein percentage (PPCT). Production in different lactations was treated as different traits. Only first, second and third lactation production traits were considered.

The original data set consisted of performance records of 200 319 Holstein cows calving between 1980 and 2005, and, pedigree information of 885 567 animals. All data were obtained from the Integrated Registration and Genetic Information System (Intergis) of South Africa. Records with unknown birth and calving dates were deleted. Age at calving within each lactation was restricted to remove outliers, using the following ranges, as used by Mostert *et al.* (2006): 20-42 months for lactation 1, 30-54 months for lactation 2 and 40-67 months for lactation 3. Records with CI less than 300 days or greater than 600 days were also discarded. The data were edited further to remove lactation records that were incomplete and unusable for genetic evaluations, according to the conditions used by the National Dairy Cattle Performance Testing Scheme of South Africa (National Dairy Cattle Performance Testing Scheme, 1999). Further edits were done to remove milk yields less than 1000 kg or greater than 30 000 kg; butterfat percentages less than 2% or greater than 9% and protein percentages less than 2% or greater than 6%.

Two calving seasons were defined as summer (October – March) and winter (April – September) (Mostert *et al.*, 2006). Each CI and AFC observation was assigned to a calving herd-year-season contemporary group. Cows with production records were also assigned to herd-year-season of calving contemporary groups. Contemporary groups with less than five animals or less than two sires were removed. Each cow had to have a first lactation record to take into account the selection of cows which made subsequent records. Finally, a random subset of data was extracted, based on contemporary group, to fit available computer capacity. A pedigree file containing all animals in the data set, going back four generations, was constructed. The pedigree file contained 97 320 cows, daughters of 4 194 sires and 60 183 dams. Table 1 shows the structure of the data set.

Descriptive statistics of all traits were computed using the Proc Means procedure of the Statistical Analysis System (SAS, 1998). An Analysis of Variance was carried out using the General Linear Model (GLM) procedure of SAS (1998) to determine non-genetic factors affecting each trait, in order to determine the effects that should be included in the models for variance component estimation. The effects tested were herd-year-season of calving, linear and quadratic effects of calving age, and previous CI (only fitted for second and third lactation production traits). All these effects were significant ($P < 0.05$) except the quadratic effect of calving age, which was not significant for all production traits.

The Variance Component Estimation-Restricted Maximum Likelihood (VCE-REML) Version 5.0 programs of Groeneveld *et al.* (2003) were used to estimate variance and covariance components. A multitrait analysis of all 17 traits simultaneously was not computationally feasible. For this reason, a series of four-trait analyses (one fertility trait and three production traits) was performed. The four-trait animal model equations 1 and 2, in matrix notation, were used respectively for the analysis of AFC and CI, each with three production traits.

Table 1 Structure and descriptive statistics of data

Trait	No of Records	HYS	Sires	Herds	Min	Max	Mean	σ_p
AFC (months)	9336	760	1 263	470	20	41	28	4
CI (days)	16 183	899	2 029	604	300	600	396	58
1 st lactation								
MLKY1 (kg)	20 618	1 269	1 862	536	1 126	15 572	6 802	1 969
BFATY1 (kg)	20 593	1 269	1 861	536	40	580	239	72
PROTY1 (kg)	20 592	1 269	1 862	536	40	497	216	61
BFPCT1 (%)	20 593	1 269	1 861	536	2.00	5.77	3.53	0.39
PPCT1 (%)	20 593	1 269	1 862	536	2.46	4.54	3.18	0.20
2 nd lactation								
MLKY2 (kg)	8 704	672	1 214	248	1 876	17 635	8 095	2 230
BFATY2 (kg)	8 661	671	1 209	248	67	691	282	83
PROTY2 (kg)	8 653	671	1 208	248	60	540	255	69
BFPCT2 (%)	8 661	671	1 209	248	2.03	5.42	3.51	0.40
PPCT2 (%)	8 653	671	1 208	248	2.58	4.06	3.18	0.20
3 rd lactation								
MLKY3 (kg)	4 112	396	829	154	2 025	28 391	8 695	2 234
BFATY3 (kg)	3 948	390	798	149	72	797	302	72
PROTY3 (kg)	3 942	390	798	151	73	567	271	73
BFPCT3 (%)	3 948	390	798	149	2.01	5.50	3.49	0.39
PPCT3 (%)	3 942	390	798	151	2.51	4.34	3.15	0.19

AFC - age at first calving; CI - calving interval; MLKY - lactation milk yield; BFATY - butterfat yield; PROTY - protein yield, BFPCT - butterfat percentage; PPCT - protein percentage

HYS - herd-year-season, Min - minimum value, Max - maximum value, σ_p - standard deviation

$$\begin{bmatrix} y_1 \\ y_2 \\ y_3 \\ y_4 \end{bmatrix} = \begin{bmatrix} X_1 b_1 + Z_1 u_1 \\ X_2 b_2 + Z_2 u_2 \\ X_3 b_3 + Z_3 u_3 \\ X_4 b_4 + Z_4 u_4 \end{bmatrix} + \begin{bmatrix} e_1 \\ e_2 \\ e_3 \\ e_4 \end{bmatrix} \quad [1]$$

$$\begin{bmatrix} y_1 \\ y_2 \\ y_3 \\ y_4 \end{bmatrix} = \begin{bmatrix} X_1 b_1 + Z_1 u_1 \\ X_2 b_2 + Z_2 u_2 \\ X_3 b_3 + Z_3 u_3 \\ X_4 b_4 + Z_4 u_4 + Wp \end{bmatrix} + \begin{bmatrix} e_1 \\ e_2 \\ e_3 \\ e_4 \end{bmatrix} \quad [2]$$

where: y_1 , y_2 and y_3 are vectors of observations for three production traits; y_4 in Equation 1 is a vector of observations for AFC and y_4 in Equation 2 is a vector of observations for CI; X_1 , X_2 , X_3 are incidence matrices relating observations on production traits to fixed environmental effects (calving HYS, linear calving age and CI); X_4 in equation 1 is an incidence matrix relating AFC observations to fixed environmental effects (calving HYS); X_4 in equation 2 is an incidence matrix relating CI observations to fixed environmental effects (calving HYS, linear and quadratic calving age); b_1 , b_2 , b_3 and b_4 are vectors of fixed effects; Z_1 , Z_2 , Z_3 and Z_4 are incidence matrices relating observations to random animal effects; u_1 , u_2 , u_3 and u_4 are vectors of additive genetic effects of the animal; W is the incidence matrix relating observations

to permanent animal environmental effects; p is the vector of random permanent animal environmental effects; e_1, e_2, e_3 and e_4 are vectors of random residual effects.

Results

Table 1 provides a summary of the data and descriptive statistics of all traits in the data set used for parameter estimation. The average CI and AFC were respectively 396 days and 28 months. Phenotypic standard deviations were 58 days and four months respectively for CI and AFC. The means for yield traits ranged from 6802 kg, 239 kg and 216 kg, for MLKY1, BFATY1 and PROTY1, respectively, to 8695 kg, 302 kg and 271 kg for MLKY3, BFATY3 and PROTY3, respectively. Phenotypic standard deviations were in the range 1969 kg, 72 kg and 61 kg for MLKY1, BFATY1 and PROTY1, respectively, to 2234 kg, 72 kg and 73 kg for MLKY3, BFATY3 and PROTY3, respectively. The phenotypic means for percentage traits ranged from 3.49% and 3.15% for BFPCT3 and PPCT3 respectively, to 3.53% and 3.18% for BFPCT1 and PPCT1.

Heritability estimates for all traits are given in Table 2. Age at first calving had a moderate heritability (0.24) while the estimate for CI was low (0.03). Production traits were moderate to highly heritable, with estimates ranging from 0.19 (BFATY2) to 0.65 (PPCT1). Estimates were highest in first lactation for all traits but there was no observable trend across lactations.

Table 2 Estimates of heritabilities for fertility and first, second and third lactation production traits

Trait	Heritability \pm s.e.
AFC	0.24 \pm 0.02
CI	0.03 \pm 0.01
1 st lactation	
MLKY1	0.33 \pm 0.02
BFATY1	0.24 \pm 0.02
PROTY1	0.28 \pm 0.02
BFPCT1	0.46 \pm 0.02
PPCT1	0.65 \pm 0.02
2 nd lactation	
MLKY2	0.25 \pm 0.02
BFATY2	0.19 \pm 0.02
PROTY2	0.24 \pm 0.02
BFPCT2	0.35 \pm 0.03
PPCT2	0.55 \pm 0.02
3 rd lactation	
MLKY3	0.25 \pm 0.03
BFATY3	0.22 \pm 0.03
PROTY3	0.26 \pm 0.03
BFPCT3	0.36 \pm 0.04
PPCT3	0.47 \pm 0.03

AFC - age at first calving; CI - calving interval; MLKY - lactation milk yield; BFATY - butterfat yield; PROTY - protein yield, BFPCT - butterfat percentage; PPCT - protein percentage

Estimates of genetic correlations between AFC and milk production traits are given in Table 3. All genetic correlations between AFC and yield traits were negative, while correlations between AFC and percentage traits were essentially zero, except that with third lactation BFPCT (0.14). Genetic correlations between AFC and yield traits decreased with increase in parity. Estimates of genetic correlations between CI and milk production traits are given in Table 3. Positive and moderate to high genetic correlations were observed between CI and yield traits, ranging from 0.37 with second lactation MLKY to 0.69 with first lactation MLKY. Genetic correlations between yield traits and CI were highest in first parity and lowest in second parity. Correlations between CI and percentage traits were low and close to zero for lactations 2 and 3.

Table 3 Estimates of genetic correlations fertility and first, second and third lactation production traits

Traits	Age at first calving	Calving interval
1 st lactation production		
MLKY1	-0.43 ± 0.05	0.69 ± 0.06
BFATY1	-0.50 ± 0.05	0.66 ± 0.06
PROTY1	-0.49 ± 0.05	0.68 ± 0.06
BFPCT1	0.03 ± 0.02	-0.17 ± 0.07
PPCT1	0.03 ± 0.02	-0.16 ± 0.06
2 nd lactation production		
MLKY2	-0.35 ± 0.05	0.37 ± 0.10
BFATY2	-0.35 ± 0.06	0.44 ± 0.10
PROTY2	-0.39 ± 0.04	0.41 ± 0.10
BFPCT2	-0.02 ± 0.07	0.05 ± 0.01
PPCT2	-0.05 ± 0.05	0.03 ± 0.01
3 rd lactation production		
MLKY3	-0.29 ± 0.08	0.52 ± 0.10
BFATY3	-0.17 ± 0.09	0.50 ± 0.10
PROTY3	-0.25 ± 0.08	0.50 ± 0.10
BFPCT3	0.14 ± 0.03	0.06 ± 0.08
PPCT3	0.08 ± 0.03	-0.03 ± 0.04

MLKY - lactation milk yield; BFATY - butterfat yield; PROTY - protein yield,
BFPCT - butterfat percentage; PPCT - protein percentage

Discussion

The primary objective of this study was to estimate genetic parameters among female fertility and production traits, to determine the effect that selection on production *per se* may have on female fertility. Yields in different lactations were treated as different traits in order to determine whether correlations were constant across lactations.

The heritability of AFC was higher than estimates obtained for Holstein-Friesian cattle in the USA (Seykora & MacDaniel, 1983). Rege (1991) and Ojango & Pollot (2001), however, reported higher heritability estimates in Kenyan Holstein-Friesian cattle. The discrepancies could be due to differences in genetic variation among the populations, differences in statistical models used for analysis or varying reactions of the same breed to different environmental conditions. Heritability estimates for production traits were comparable to those obtained previously for the South African Holstein population (Tesfa, 2002) and

generally within the range reported in the literature (Pryce *et al.*, 1998; Lobo *et al.*, 2000; Haile-Mariam, 2003; Kadarmideen, 2004). The moderate heritabilities for AFC and production traits indicate that there is potential for improvement of these traits through selection. The low heritability estimate for CI is in agreement with many other studies on Holstein-Friesian cattle (Hoekstra *et al.*, 1994; Veerkamp *et al.*, 2001; Olori *et al.*, 2002; Wall *et al.*, 2003; 2004). However, higher estimates were reported for Holstein-Friesian cattle in Kenya (Rege, 1991; Ojango & Pollot, 2001) and Florida, USA (Campos *et al.*, 1994).

The low heritability estimate for CI indicates that significant improvement through selection is attainable only through enhanced accuracy of selection by, for example, incorporating information on correlated traits. Traits such as body condition score, certain linear type traits, milk progesterone and milk urea nitrogen have been shown to be genetically correlated with fertility (Dadati *et al.*, 1986; Darwash *et al.*, 1999; Melendez *et al.*, 2000; Pryce *et al.*, 2000; Royal *et al.*, 2000; 2002; Dechow *et al.*, 2001; Godden *et al.*, 2001; Veerkamp *et al.*, 2001; Gutierrez *et al.*, 2002; Berry *et al.*, 2003; Wall *et al.*, 2003; 2005; Haile-Mariam *et al.*, 2004; Kadarmideen, 2004).

Estimates of genetic correlations between fertility traits (AFC and CI) and milk production traits from most previous studies were only based on first lactation production (Rege, 1991; Pryce *et al.*, 2000; Ojango & Pollot, 2001; Campos *et al.*, 1994; Hoekstra *et al.*, 1994). Genetic correlations observed between AFC and milk yield were higher and more negative than estimates observed in Kenyan Holstein-Friesian cattle by Rege (1991). Across all lactations, genetic correlations between AFC and milk yield were, however, smaller than estimates obtained by Ojango & Pollot (2001), also in Holstein-Friesian cattle in Kenya. Correlations were highest in first parity and lowest in third parity, indicating a strong relationship between these traits in early parities. The negative association between AFC and yield traits indicates a favourable effect of yield traits on age at first calving. Thus, selection for increased yield may result in a correlated improvement in AFC. The stress of increased yield may, however, overcome the genetic merit for improved fertility, especially *post-partum* re-conception (Rege, 1991). Correlations between AFC and percentage traits were mostly close to zero, except that between AFC and third lactation butterfat (0.14), which was similar to that reported by Rege (1991). This indicates that selection for butterfat percentage is detrimental to AFC.

The genetic correlations between CI and yield traits were moderate to high and positive, in agreement with most literature estimates for Holstein-Friesian cattle (Hoekstra *et al.*, 1994; Pryce *et al.*, 1997; 1998; 2000; Ojango & Pollot, 2001; Veerkamp *et al.*, 2001). Smaller estimates were, however, reported for Holstein-Friesian cattle in Kenya by Rege (1991) and in the United Kingdom (Wall *et al.*, 2003). Genetic correlations between CI and yield traits were highest in first parity, indicating that *post-partum* fertility is mostly associated with first lactation yield. A positive genetic correlation between CI and yield indicates that animals with high genetic merit for milk, butterfat and protein yield would tend to have longer calving intervals, indicating antagonism between CI and yield traits. Similar antagonistic relationships have been reported in studies between most fertility traits and production (Hoekstra *et al.*, 1994; Veerkamp *et al.*, 2001; Berry *et al.*, 2003; Wall *et al.*, 2003). Due to this fact, intense selection for yield, without regard to fertility has led to deterioration in reproductive performance in most countries (Van Arendonk *et al.*, 1989; Frick & Lindhe, 1991; Bagnato & Oltenacu, 1994; Campos *et al.*, 1994; Hoekstra *et al.*, 1994; De Jong, 1998; Pryce *et al.*, 1997; Ojango & Pollot, 2001; Kadarmideen, 2004; VanRaden *et al.*, 2004). The genetic correlations between CI and protein and butterfat percentage traits were low for lactation 1 and close to zero for lactations 2 and 3, suggesting that there is low genetic association between CI and percentage traits.

Conclusions

The heritability estimate for age at first calving and production traits indicate that these traits can be improved genetically through selection in the South African Holstein population. Selection on CI is, however, likely to achieve slow progress, due to its low heritability. There is a need to enhance the accuracy of selection on CI by increasing information used for its evaluation.

Findings indicate favourable genetic relationships between age at first calving and yield in South African Holsteins. However, unfavourable genetic associations were found between calving interval and yield traits. A correlated decline in *post partum* cow fertility (increased CI) is likely to result from selection on yield, without regard to fertility. Thus, female fertility should be included in the breeding objective for South African Holstein cattle.

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