

GENETIC AND PHENOTYPIC CORRELATIONS OF AGE AT FIRST CALVING WITH SOME CALF-HOOD FITNESS AND HEALTH TRAITS IN HOLSTEIN HEIFERS

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SUMMARY

Objectives of this study were to estimate the effects of calf-hood fitness and health traits on age at first calving (AFC). Data of 2230 US Holstein heifers were collected from three farms located in Minnesota from 2003 to 2011 were used. AFC, calculated as a difference between birth date and date of first calving ranged from 571 to 1059 days ($\bar{x} = 716.66 \pm 47.4$). Multiple trait animal models were applied using REMLF90 software. Fixed effects considered as revealed in; a previous nutritional experiment applied on almost half of these animals ($n = 1226$), ($P < 0.01$), and interaction between month and year of birth, while covariate effects were serum protein upon arriving to facility (SP), ($P < 0.01$). Fitness traits effects included arrival to facility body weight (ABW), weaning weight (W56), weaning hip height (HH56), adjusted weight to 200d (W200), and adjusted hip height to 200d (HH200). Health traits recorded were scours (Sc) and respiratory disease (RD). Health traits were recorded as categorical traits according to incidence of occurrence (0: non occurrence; 1: occurrence once; and 2: occurrence twice or more). Heritability estimate of AFC was (0.07). Genetic correlations between calf-hood fitness traits and AFC were highly significant (0.45, 0.38, 0.34, 0.75, and 0.33) with ABW, W56, HH56, W200, and HH200, respectively. Meanwhile, genetic correlations between AFC and both health traits (Sc and RD) were (-0.19 and -0.01), respectively. Therefore, it is possible to build a selection index comprising calf-hood's fitness and health performance and AFC.

Keywords: age at first calving, fitness traits, health traits, heritability, genetic correlation

INTRODUCTION

Age at first calving is defined as the number of days from birth to day of first calving. At this stage, replacement heifers are not generating any income. Instead, this rearing period requires considerable capital expenditures including feed, housing, and veterinary expenses that had been estimated by Heinrichs (1993) as 15 to 20% of the total expenses related to milk production. Cooke *et al.* (2012) showed that rearing replacement heifers is considered a costly process where early AFC can reduce this cost.

Tozer and Heinrichs (2001) showed that reducing AFC from 25 to 24 or 21 months decreased replacement costs by 4.3% or 18%, respectively. Meanwhile, increasing AFC to 29 months led to increase cost by 14%, also led to keep more replacement heifers in herds to maintain herd size. Meanwhile, Hoffman *et al.* (1992), reported that acceptable milk yield and levels of dystocia could be achieved in modern Holstein heifers when calving at ages < 22 mo due to genetic selection and advances in nutritional management.

Previous studies on heritability estimates of AFC in different breeds are shown in Table (1).

Table 1. Estimates of heritability for AFC in different breeds

Breed	Estimate	Statistical models	References
Boran cattle in Kenya	0.04±0.06	Univariate animal models	Wasike <i>et al.</i> (2009)
Iranian Holstein	0.014±0.005	NA*	Chookani <i>et al.</i> (2010)
Angus Orejinegro Zebu	0.15±0.13	Multiple trait mixed model	Vergara <i>et al.</i> (2009)
Romosinuano	0.16±0.09	Means of linear models	Suárez <i>et al.</i> (2006)
South African cattle	0.26 ±0.02	Multiple trait animal model	Makgahlela <i>et al.</i> (2008)
Iranian Holstein	0.086	Animal model	Nilforooshan and Edriss (2004)
Iranian Holstein	0.19±0.005	Single and two trait animal model	Arough <i>et al.</i> (2011)

*: NA: Not available.

Ettema and Santos (2004) evaluated the effects of altering age at breeding to affect a change in AFC by using Holstein heifers, their results showed that medium and high AFC increase rearing cost than low

AFC by 40.34\$ and 107.89\$ for each heifer, respectively.

In the past, growth has always been defined as weight gain per day. Other body size criteria such as

wither height, body condition score, body length, and pelvic area are now available to aid growth definitions. These measurements should be used because they are related to lactation performance and animal health than body weight. Hoffman *et al.* (1997) used three ranges (lower, average, and upper) for weight, wither height, body length, pelvic area, and body condition score (BCS) as a criteria for a recommended optimum body size of Holstein replacement heifers at first calving.

One of the aims of this study was to estimate genetic correlations between health traits and AFC. Unfortunately, there is no direct selection for health traits in calves due to: 1) difficult and expensive health data recording; 2) needs a large population to follow up and diagnosis; 3) health traits have low heritability, Heringstad (2010). Because of the above reasons, health traits have been the last to be incorporated into genetic evaluation programs for dairy calves. Steine *et al.* (2008) reported that only in Nordic countries, direct selection for improved disease resistance had been carried out for more than 30 years.

In 2007, a national dairy cattle health and disease data management system was initiated in Canada. The main objective was to provide dairy producers and veterinarians with information on herd management and to establish a national genetic evaluation system for genetic selection for disease resistance, Koeck *et al.* (2012). Calving survival traits had gained attention recently especially for beef breeders. These traits are assumed to have underlying continuous genetic and environmental influences, but the problem is the low heritability estimates of these traits, Guerra *et al.* (2006). USDA (2007) performed a comprehensive study to document calf losses in North America and found that 15.9% of calves die before weaning, 8.1% of this percentage resulted from events that occur during calving. These losses are classified as stillbirths. The remaining deaths (7.8%) were associated with health problems caused by pathogens acquired after birth, most commonly manifested as diarrhea and respiratory problems leading to calf death. Heringstad *et al.* (2008) reported that there could be a possible association between calf diseases and other economic traits recorded later in cow's life, such as growth, fertility, and production, therefore from a breeding perspective, data on calf-hood diseases may be valuable.

Morrison *et al.* (2013) reported that calves born with no immunity, therefore feeding colostrum is important since its value in disease management has long been known. They also reported that calves with poor colostrum intake and inadequate immunity in early life were found to cause a 40% increase in treatment costs on farm. Most common parameter used under field conditions to measure positive or failure of passive transfer of immunoglobulins (Ig) is serum total protein (SP) since it is highly correlated with immunoglobulins especially IgG levels.

The aim of the present study is to focus on calf-hood fitness and health traits on their effects on AFC and to investigate if it is possible to build selection indices to select calves with a reduction in AFC. Unfortunately, studies taking calf-hood traits and following dairy calves through calving, their first lactation, and their complete lifetime production are few.

MATERIALS AND METHODS

Age at first calving was calculated as the difference between birth date and date of first calving in days and was available on 2230 Holstein heifer calves collected from three commercial dairy farms picked up twice weekly at two to four days of age and remained at the University of Minnesota, Southern Research and Outreach Center (SROC) Calf and Heifer Research and Extension facility in Waseca, Minnesota up to six to seven months of age.

During winter months, calf blankets are used at pick-up and remain on the calves at the discretion of (SROC) staff until they adjust to their new environment. In the nursery phase, calves are housed in one of two 60 m x 9 m curtain side-wall naturally ventilated calf barns. Each barn contains two 27 m x 9 m rooms with 40 individual pens (approx nine sqm /calf) within each room. A six m x nine m mixing and feed storage area is centrally located in each barn. The rooms are managed as an all-in, all out system. All pen panels are removed and power washed between calf groups. All bedding material is removed and the remaining front gates and rear panel holders are also power washed. In winter, chopped straw is used for bedding calf pens while sawdust is used in summer months.

Upon arrival, calves are weighed, and two jugular blood samples drawn. One sample is used to check total serum proteins using a refractometer (scale of serum protein). A second sample for whole blood analyses by an outside laboratory to identify persistently infected of Bovine virus diarrhea (BVD) calves.

Nearly half of calves ($n = 1226$) used in this study were involved in a nutritional experiment, therefore this trait was added as a fixed effect in the model in order to eliminate this effect on studied traits.

Analyses of variance using GLM procedures (SAS, 2004) were conducted to test the fixed effects of farm, year of birth, month of birth, season of birth, a previous nutritional experiment applied on nearly half of these animals ($n = 1226$), the interactions between month of birth and year of birth, year and season, and year and farm, revealed no significant effect ($P > 0.10$) of year of birth, month of birth, season of birth and interactions between year X season, and year X farm, therefore they were deleted from the final model in order to calculate the least squares means for each level of each factor.

Animal mixed models were applied by using REMLf90 software package, Misztal *et al.* (2002) to estimate heritability of AFC, phenotypic, genetic and

residual correlations between calf-hood traits and AFC. The following model was applied to fit the data:

$$Y_{ijklmn} = h_i + mp_{jk} + t_l + s_m + a_n + e_{ijklmno}$$

where:

- $Y_{ijklmno}$ = observation of studied traits;
- h_i = fixed effect of i^{th} herd $i = (1, \dots, 3)$;
- mp_{jk} = interaction between j^{th} month of birth and k^{th} year of birth;
- t_l = fixed effect of treatments within nutritional trails¹;
- s_m = covariate effect of serum protein;
- a_n = random animal effect;
- $e_{ijklmno}$ = random residual effect associated with the observation $ijklmno$.

1: this effect was excluded when estimating ABW.

In matrix notation form, the multiple trait models can be described as:

$$Y = Xb + Zu + e,$$

where:

- Y = vector of observations of studied traits;
- X = incidence matrix relating the observations to the fixed effects;
- b = vector of fixed effects;
- Z = incidence matrix relating the observations to the random effects;
- u = vector of random effects;
- e = vector of residual effects;

The expectations and assumed variances are $E(y) = Xb$; $E(u) = E(e) = 0$; $V(u) = G$; $V(e) = R$; $cov(u, e) = 0$; and $V(Y) = ZGZ' + R$.

Table 2. Structure and descriptive statistics for studied traits

Observations ^a	Mean	SD	Min.	Max.
SP(g/dl)	5.4	0.77	2.6	9
ABW (Kg)	39.4	5.0	20.86	58.96
W56 (Kg)	73.8	9.2	43.9	104.8
W200 (Kg)	206.6	21.2	68.04	288
HH56 (cm)	89.6	3.3	74.93	98.4
HH200 (cm)	114.4	3.6	100.33	127
AFC (d)	716.7	47.4	571	1059

a:SP: Serum Protein; ABW: Arrival body weight; W56: Weight at 56d; W200:Weight after standardized to 200d; HH56: Hip Height at 56d; HH200: Hip Height standardized to 200d; Sc: scours; RD: respiratory disease; AFC: age at first calving.

Table 3. Incidence of occurrence for health traits

Health traits ^a	Incidence occurrence (%)		
	0	1	2
Sc	48.83	49.87	1.30
RD	92.6	6.91	0.13

a:Sc: scours; RD: respiratory disease

As shown in Table (2), mean of SP was 5.4 with a range of 2.6 to 9 g/dl, Chester-Jones and Broadwater (2009) showed that if calves have received enough high quality colostrum, serum total protein will be 5.5 grams per deciliter (g/dl) or greater, if the value falls between 5.0 and 5.5 g/dl, there is a marginal risk for mortality and morbidity, if the value is less than 5.0 g/dl the calf will be at high risk for health problems. Villarroel *et al.* (2013) reported that maximum concentration of SP is when calves reach 2 - 3 days of age. After that, concentrations decreased

The convergence criteria used in this analysis was the variance of simplex values (-2 log likelihood) of $< 10^{-6}$ and the value of (-2 log likelihood) did not change from the previous solution by at least two decimal points. To ensure global convergence, programs were restarted with previous solutions at local convergence until convergence occurred at the same maximum. Overall, number of iterations required to reach convergence could be affected by the number of animals, number of random effects in the model and number of traits studied, (El-Arian *et al.*, 2003).

The animal pedigree file was generated by tracing the pedigrees of cows with data 11 generations back and contained the relationship of 26.553 animals.

RESULTS AND DISCUSSION

Mean, standard deviation (SD), and range of fitness traits included calf weight once arrived to SROC facility (ABW) since these animals were picked twice weekly from origin farms, weaning weight (W56) at 56d, weaning hip height (HH56), adjusted weight to 200d (W200), and adjusted hip height to 200d (HH200) with the reproductive trait: age at first calving (AFC) are shown in Table (2).

Percentage of incidence of occurrence of diseases scours and respiratory disease are recorded in Table (3).

by approx. 0.07 g/ dL, this statement indicates that these estimates were at the maximum concentration of SP since these estimates were taken at the same age (2-3days).

Mean of W56, and W200 were 73.8, and 206.6 kg, respectively. Heinrichs and Hargrove (1986) working on 5723 Holstein heifers collected from 163 herds found the mean weight of 2 mo and 6 mo as 82.1, and 168.6 kg, respectively. Meanwhile, mean estimates for HH56, and HH200 were 89.6, and 114.4 cm respectively. Results from Heinrichs and

Hargrove (1986) showed 85.6 and 101.1 for height at two and six months, respectively. Difference in values shown between studied traits and Heinrichs and Hargrove (1986) traits could be due to different days on taking these values in different studies.

Mean age at first calving was 716.7 days (24 month). Makgahlela *et al.* (2008) showed that the mean AFC in South African Holstein found it as 840 days (28 month), also several authors (Arough *et al.*, 2011 and Chookani *et al.*, 2010) found the mean of AFC for Iranian Holstein as 811.1 days (26.6 month). Mean of AFC according to year are shown in figure (1). It is obvious that the protocol in (SROC) University facility is to low AFC in order to reduce rearing costs due to decreased feed, labor and housing costs.

Incidence of occurrence of Sc for years from 2003 to 2011 combined was 51.17% (as shown in Table 3). Stoltenow and Vincent, (2003) showed that

the first three weeks of the calf life is important because any infection on that age resulted in some level of scours due to the fact that its gut is still so immature during that period and considered the weakest point of the calf's system. The cause could be a single agent such as salmonella. But if a variety of agents are present, the diagnosis of the causal pathogen cannot be accurately made.

Incidence of occurrence for RD (both 1 and 2) was 7.04%, these estimate was lower than Snowder *et al.* (2005) working on 18,112 records from 9 different beef breeds collected from 1987 – 2001 reported that incidence of RD exceeded 20% in early years (1987 -1991), while it lowered to 14% in late years (1992 – 2001). They showed that the reason for this was; 1) modifying vaccines with higher effective in later years and; 2) changing castration method from surgical to mechanical.

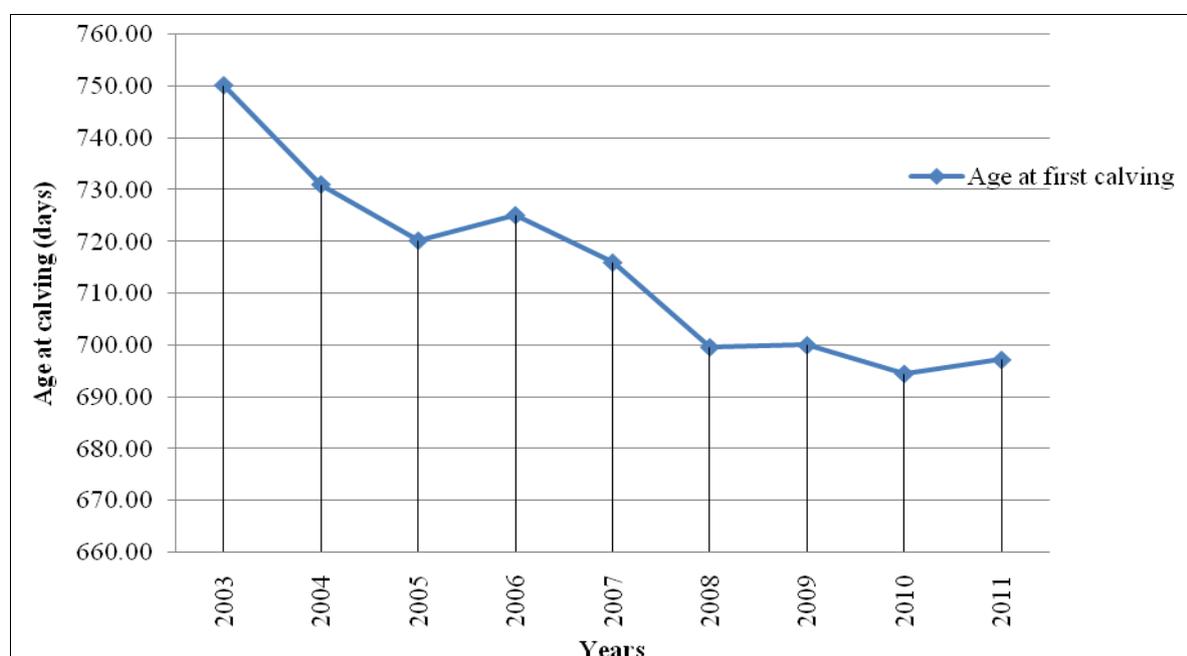


Fig. 1. Mean estimate of age at first calving according to year

Heritability estimate for AFC in this study was (0.07). This estimate was close to Wasike *et al.*, (2009) for Boran cattle in Kenya (0.04 ± 0.06), Chookani *et al.* (2010) for Iranian Holstein (0.014 ± 0.005), Nilforooshan and Edris (2004) in Iranian Holstein (0.086). Meanwhile, this estimate was lower than (0.15 ± 0.13) obtained by Vergara *et al.* (2009) for Angus Blanco Orejinegro Zebu cattle in Colombia, (0.16 ± 0.09 ; Suárez *et al.*, 2006) on Romosinuano in Colombia cattle, (0.19 ± 0.005 ; Arough *et al.*, 2011) on Iranian Holstein. On the other hand, Makgahlela *et al.* (2008) reported a medium heritability estimate of AFC on South African cattle as (0.26 ± 0.02).

Arough *et al.* (2011) reported that there could be several factors for the difference in heritability estimate such as: breed of animal, management system, environmental factors, size and structure of data, model of analyses, and statistical methods

employed. Several authors (Kadernideen 2004, Makgahlela *et al.*, 2008) reported that, due to the low heritability for AFC, selection for improving this trait in dairy cattle would not worthwhile.

Considering AFC as a fertility trait, De Jong (1998) found high additive genetic variation in fertility traits. Accordingly it could be possible to improve these traits genetically through selection, and this could be achieved by increasing the amount of information used in the genetic evaluation (e.g., using offspring information). Incorporation of traditional measures of fertility and all these correlated traits, directly and indirectly, could be used to improve the accuracy of genetic predictions for fertility traits (Makgahlela *et al.*, 2008).

It is stated that the cow become profitable when entering the second lactation, therefore several farmers considers calf-hood stage and the first lactation stage as considerable capital expenditures.

Lin *et al.* (1988), and Van Amburgh *et al.* (1998) reported that milk yield in second and greater lactations has consistently been unaffected by reduced AFC, therefore, it will be beneficial lowering AFC.

Highly significant ($P < 0.01$) genetic and phenotypic correlations between calf-hood fitness traits and AFC ranged from 0.33 to 0.75 as shown in Table (4). These estimates show a positive genetic relationship therefore, adding these traits in the selection indices will improve AFC genetically.

Table 4. Genetic correlations between calf-hood traits and AFC

Calf-hood traits ^a	AFC		
	Genetic correlation	Residual correlation	Phenotypic correlation
Fitness traits			
ABW	0.45**	-0.07*	0.38**
W56	0.38**	-0.04	0.34**
W200	0.75**	-0.13**	0.62**
HH56	0.34**	-0.04	0.30**
HH200	0.33**	-0.05	0.28**
Health traits			
Sc	-0.19**	0.00001	-0.19**
RD	-0.01	0.02	-0.03

a: ABW: Arrival body weight; W56: Weight at 56d; W200: Weight after standardized to 200d; HH56: Hip Height at 56d; HH200: Hip Height standardized to 200d; Sc: scours; RD: respiratory disease. *: significant at 0.05; **: highly significant at 0.01.

Negative genetic and phenotypic correlations between health traits and AFC as shown in Table (4) indicate that disease occurrence of both scours and respiratory illness at calf-hood stage increase AFC. This result agrees with Heinrichs *et al.* (2005) who reported that health of calves during the first 4 mo of life and antibiotic treatment for scours or pneumonia increase AFC.

CONCLUSION

Healthy, productive herd replacements are the result of good management before the calves are conceived and continue to the time when they enter the milking herd. Accurate records must be kept so a sound breeding program can be followed. The total process ensures that genetically superior animals will enter the herd. Once calves are born, they should be provided with a healthy environment including proper facilities, water, high quality feeds, daily observations, and health care. This kind of attention should extend beyond the baby calves. A carefully managed, well-planned facility enables dairy farmers to efficiently use their time and labor while rearing healthy replacements that freshen at 24 months of age.

Due to the low heritability estimate of AFC reported in this study (0.07) it seems that this trait is more controlled by environmental and management practices rather than genetically practices, it will be beneficial to add calf-hood fitness and Sc traits (according to their high significance) to the selection indices to reach the target AFC according to each farm strategies.

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المكافئ الوراثي والارتباطات الوراثية للعمر عند أول ولادة كتأثير بموائمة العجول الرضيعة والحالة الصحية للهولستين

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أهداف هذه الدراسة كانت لتقدير تأثيرات موائمة العجول الرضيعة والحالة الصحية على العمر عند أول ولادة واستكشاف اذا كانت صفات العجول الرضيعة يمكن أن تؤثر على العمر عند أول ولادة. البيانات المتوفرة على 2230 عجلة هولستين أمريكي تم تجميعها من ثلاث مزارع موجودة في ولاية ميناسوتا في الفترة من 2003 إلى 2011.

حسب العمر عند أول ولادة كفرق بين تاريخ الميلاد وتاريخ أول ولادة بالايام والتي تراوحت من 571 إلى 1059 يوم (بمتوسط قدرة 716.66 ± 1.00). تم تطبيق نموذج الحيوان متعدد الصفات باستخدام برنامج REMLf90. العوامل المصحح لها كانت تأثير المزرعة، تجربة تغذية سابقة والتي طبقت على ما يقرب من نصف الحيوانات (N = 1226) والتداخل بين الشهر وسنة الميلاد بينما تم ادراج بروتين السيرم بمجرد الوصول الى المزرعة كتأثير تغاير.

الموائمة شملت الوزن عند الوصول للمزرعة (ABW)، وزن الفطام (W56)، ارتفاع الخصر عند الفطام (HH56)، الوزن المعدل ل 200 يوم (W200) وارتفاع الخصر المعدل ل 200 يوم (HH 200). الصفات الصحية المسجلة كانت الاسهال (Sc) وامراض التنفس (RD). هذه الصفات الصحية كانت قد سجلت كصفات تدريجية طبقاً لمعدلات الحدوث الأصابة (صفر: غير موجود، واحد: موجود مره واحده، اثنين: موجود مرتين أو أكثر). قدر المكافئ الوراثي للعمر عند أول ولادة ب (0.07). الارتباطات الوراثية بين صفات الموائمة والعمر عند أول ولادة كانت (0.45، 0.38، 0.34، 0.75 و 0.33) لكلاً من وزن الجسم عند الوصول للمزرعة، وزن الفطام، ارتفاع الخصر عند الفطام، الوزن المعدل ل 200 يوم وارتفاع الخصر المعدل ل 200 يوم على التوالي. بينما كان الارتباط الوراثي بين العمر عند أول ولادة وكلاً من صفتي الحالة الصحية (الاسهال والامراض التنفسية) (-0.19 و -0.01) على التوالي. طبقاً لهذه النتائج فأنه يمكن بناء دليل أنتخاب يشمل موائمة العجول الرضيعة والاداء الصحي والعمر عند أول ولادة.