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RESEARCH ARTICLE

Estimation of Genetic Parameters using Animal Model for Some Performance Traits of Nili-Ravi Buffaloes

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ABSTRACT

The objective of this study was to estimate the genetic parameters of productive traits of Nili-Ravi buffaloes including milk yield (MY), lactation length (LL) and dry period (DP) for the identification of superior animals. For this purpose, data on 9003 lactation records of Nili-Ravi buffaloes from four institutional herds in Punjab, Pakistan were used for the present study. Model incorporating all known relationships was used for estimating variance components by ASREML and DFREML programs. The heritability evaluations for 305-day milk yield (305-day MY), total milk yield (TMY) and LL ranged from 9 to 23, 10 to 18 and 7 to 18 percent, for different parities. The overall heritability for 305-day MY, TMY, LL and DP, were 9, 10, 6 and 3 percent, respectively. The repeatability estimates for the 305-day MY, TMY, LL and DP were 35, 36, 22 and 9 percent, respectively. The low estimations for genetic parameters for all the production traits proposed that most of the observed variant in these traits was due to environmental settings or non-additive genetic effects. However high genetic variability indicates that there is an ample opening for the advancement of these productive traits while including in the future breeding programs.

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INTRODUCTION

The Nili-Ravi buffalo originated during the sixties under a major policy decision of Livestock and Dairy Development Department of Government of Punjab considering Nili and Ravi as one instead of two independent breeds (Shah, 1991). The home tract of "Nili" breed is the Valley of Sutlej River, particularly Vehari, Pakpattan and Sahiwal Districts. It is one of the finest buffalo breeds in the world for milk production, body conformation and temperament. The home tract of other famous breed "Ravi" lies in Sandal Bar area of Ravi river valley in Faisalabad, Okara and Sahiwal Districts. In the second half of the last century, as communication increased and grazing lands started decreasing, the two breeds (Nili and Ravi) started mixing in a way that their distinctive characteristics merged with each other (Shah, 1991). The Nili-Ravi, thus, was recognized as a separate breed. It is now most common buffalo breed of Pakistan (37.7 million heads) and has a major contribution (34 million tonnes) in total milk production in the country (GOP, 2017). The Nili-

Ravi is the most popular and most productive breed for the commercial setup. It has been exported to countries like China, Kenya, Egypt and Australia. Most of the genetic estimations of productive traits for buffaloes came from Government herds in India and Pakistan. Genetic parameters have been reported to vary considerably. Heritability (h^2) estimates for milk yield in Pakistani buffaloes (Khan et al., 1997a), estimated using a Multiple Trait Animal Model, ranged between 17-18%. Thevamanoharan et al. (2002) testified the heritability approximations for 305-day MY and TMY as 0.01 ± 0.02 and 0.10 ± 0.01 , respectively. Previously, Cady et al. (1983) reported that sire variance for MY decreased as records were more restricted for LL of >60 , >250 and the 305-day reduction was 4.3, 2.8 and 1.1%, respectively. The h^2 for three groups (lactation length of >60 , >250 and 305-day) was 25.4, 17.9 and 7.1%, respectively. Estimates of repeatability for three groups were 26.5, 24.5 and 22.9 %, respectively. Khan (2000) also reported that lactation length adjustment procedure may affect the h^2 estimate. Unadjusted lactation milk yield was reported to be 9% heritable

while adjusted milk yield was 12%. In Indian studies, h^2 for lactation milk yield had been reported between 0.08 to 0.65 (Jakhar et al., 2017). In Indian Murrah higher h^2 estimates for TLMY, 305MY were recorded as 0.392 ± 0.114 and 0.348 ± 0.118 , respectively. Egyptian studies on genetic parameters for lactation milk yield also indicated a wide variation of h^2 estimates (0.05 to 0.30). Most of these studies utilized data from a single experimental herd. Heritability estimates from field recorded herds had only been reported for Italian buffaloes. Pilla and Moili (1992) reported that 305-day milk yield had heritability of 0.27. Later, Rosati and VanVleck (2002) reported lower estimates (0.14) using large data set on buffaloes. Catillo et al. (2001) reported estimate for milk yield of 0.19 for Italian buffaloes. Heritability and repeatability estimates for other traits even vary more widely. Similarly, there is no consensus on genetic correlations among various traits. For example, milk yield and lactation length have been reported to high genetic correlation of -0.15 (Peeva, 1997) and 1.0 (Mohamed et al., 1993). Difference of population, methodology, season and adjustment for other factors such as lactation length are commonly recognized reasons (Bashir et al., 2015). Breed improvement reforms in buffalo are very limited (Moioli et al., 2000) and with the exception of Italian studies, limited population data have been used to estimate the genetic trends.

As implementation of a breeding program in buffalos, requires genetic variation and estimation of genetic parameters such as heritability and repeatability. The genetic configuration of a population can be studied by bearing in mind the relative significance of heredity and environmental aspects affecting it. Estimation of genetic parameters for different productive traits of Nili-Ravi buffaloes in Pakistan have previously been based on the intra-sire regression of daughter on dam and covariance of paternal half-sibs, with pre-adjustment of fixed effects in the model. Breed improvement program in buffalo in Pakistan are new (Khan, 2001) but their expansion to more institutional and private herds requires estimates of genetic parameters and genetic correlation between economic traits.

The existing study aimed to estimate the heritability and repeatability of different productive traits in Nili-Ravi buffaloes and to suggest the plan for the future breeding plan for genetic improvement regarding dairy buffaloes in the country.

MATERIALS AND METHODS

Production records of Nili-Ravi buffaloes were obtained from the four Government herds of Livestock and Dairy Development Department of Punjab province; namely Livestock Experiment Station, Haroonabad, (LESHA) from 1979-2000; Livestock Experiment Station, Chak

Katora, (LESCK) from 1971-2000; Livestock Experiment Station Khushab, (LESKH) from 1979-2000 and Livestock Production Research Institute, Bahadurnagar, (LPRIBN) from 1971-2000 for the present study. Production traits included 305-day milk yield (305-day MY), total milk yield (TMY), lactation length (LL) and dry period (DP).

Data on 305-day MY and TMY were based on actual weekly milk records. Lactations shorter than 60 days were excluded (5.2 %). Incomplete lactation records of buffaloes due to culling, abortion or diseases were also excluded from this study (Bashir et al., 2015).

Data was edited using the descriptive statistics the Statistical Analysis Systems and genetic parameters were estimated using Multiple Trait Animal Model. DFREML program was used to estimate genetic parameters (Meyer, 1997). Co-variance components were estimated with both uni-variate and bi-variate models. The basic model fitted was as follows:

$$Y = Xb + Zu + e$$

Where,

Y = Vector of observations,

X and Z = Design matrices for fixed and random effects, respectively,

b = Vector of fixed effects having herd-year of calving, calving season and age code (Iqbal, 1996),

u = Vector of individual animal breeding values

e = Vector of random error terms.

The bivariate model was used as under.

$$Y = (I_t \otimes X)b + (I_t \otimes Z_1)a + (I_t \otimes Z_2)p + e$$

Where, Y is a matrix of dependent variable having vectors of lifetime traits; b is vector of fixed effects; t is number of traits i.e. 2; \otimes is Kronecker product, a, p, and e are random animal, permanent environment and temporary environment effects, respectively; X, Z_1 and Z_2 are incidence matrices for vectors b, a and p. Fixed effects included 94 herd-year combinations.

When combined analysis was performed for first five parities 'u' in the model also had permanent environment effect. The assumptions for explanations of various vectors and matrices were similar to univariate and bivariate analysis (Khan, 1998, 1997).

RESULTS AND DISCUSSION

Lactation milk yield

A heritability estimate of 305-day MY from univariate animal model was 0.10 ± 0.02 for all lactation records (Table 1). The analysis of individual parities depicted increasing trend from 1st to 5th parities. The h^2 was highest (0.23 ± 0.11) for 5th parity. When combined analysis was performed for 1st five parities (0.10 ± 0.2) or all ten parities (0.09 ± 0.02), estimate was the same. Heritability estimate in the present study was to some extent lower than the earlier studies for Nili-Ravi in Pakistan where estimate ranged between 0.14 to 0.18 or

Indian Nili-Ravi (0.15 to 0.19), Bulgarian (0.12), Egyptain (0.13), Italian (0.14), Murrah (0.14) and Surti (0.17) buffaloes (Khan et al., 1997a, b; Tailor et al., 1992a; Pareek and Narang, 2014). The heritability of milk yield increased with the increase in lactation number. In Simmental and Brown Swiss cattle, Emmerling et al. (2002) reported the h^2 for 1st, 2nd and later parities were 0.20, 0.23 and 0.25, respectively. The change between first and later lactation for lactation milk yield were not large. The increase in h^2 across parities had also been reported for Ayrshire, Guernsey, Holstein and Jersey in Canada (Jorjani, 2000). Yang et al. (2005) also reported the increase in h^2 of milk yield from 1st to 6th parity. In present study, error variance did not significantly increase across the parities, which ultimately resulted in non-significantly different heritability estimates across the parities.

The repeatability estimates for combined analysis of all parities were found 0.35 and 0.36 for 305-day MY and TMY, respectively (Table 1) which were same when data were restricted to first five parities. Repeatability estimates obtained in this study were similar to the previous studies on Murrah, Nili-Ravi, Egyptian, Surti and Italian buffaloes (Khan et al., 1996; Hamed, 1994; Jamuna et al., 2015; Pilla and Moioli, 1992; Morammazi et al. 2010; Jakhar et al. 2017). These estimates suggested that culling for the milk yield could be done on a first lactation record to improve the performance of herd.

Lactation length

The h^2 estimate for the lactation length was 0.06 ± 0.017 (Table 1). Low h^2 for lactation length obtained in the present study are in conformity with the previous findings on Nili-Rani and Murrah buffaloes (Khan, 1997; Salah-ud-Din, 1989; Dutt and Taneja, 1995; Rana et al., 2002; Jakhar et al., 2017). The h^2 estimates from these studies ranged from 0.05 ± 0.13 to 0.07 ± 0.07 .

Repeatability of lactation length was 0.20 when 1st five lactations were used and 0.21 when all lactations were included in the data set (Table 1). The present repeatability estimate for lactation length is similar to previous reports (Khan, 1997; Khan and Chaudhry, 2000) in Pakistani buffaloes. A very low estimate of 0.04 for Murrah was reported by Sesana et al. (2014) but later studies (Pareek and Narang, 2014) reported higher estimates (0.17 to 0.25) for the Murrah breed. The estimates of repeatability in Egyptian buffaloes were even higher (0.21 to 0.45) as reported by Ashmawy (1991) and Ayyat et al. (1996) and in cattle by Ayalew et al. (2017).

Dry period

The h^2 estimate for dry period in present study based on 7283 lactation records was 0.03 ± 0.01 (Table 1). The h^2 estimate obtained in the present study is similar to the Pakistani (0.07 to 0.09), Egyptian (0.07 to 0.13) and Indian buffaloes (0.08 to 0.11) (Thevamanoharan et al., 2002; Aziz et al., 2001; Khan, 1997; Ayyat et al., 1996; Dutt and Taneja, 1995; Tailor et al., 1992b). The h^2

Table 1: Heritability (h^2) and repeatability (r) estimates with standard errors, additive (σ_a^2), phenotypic (σ_p^2), environmental variances and coefficient of variation (CV) for the productive traits

Traits	N	$h^2 \pm SE$	σ_a^2	σ_p^2	σ_e^2	CV	r
305-day milk yield (kg)							
Parity 1	2050	0.136±0.04	39767.98	291522.90	251754.9	32.08	
Parity 2	1797	0.160±0.05	58461.64	363730.27	305268.6	34.28	
Parity 3	1533	0.145±0.05	48835.93	335561.31	286725.4	32.36	
Parity 4	1192	0.209±0.06	66499.53	317870.19	251370.7	32.36	
Parity 5	875	0.235±0.11	77007.66	326433.25	249425.6	32.50	
upto 5th parity	7446	0.105±0.02	35501.40	337255.50	215886.3	33.34	0.359
All parities	9003	0.095±0.02	31908.85	335519.57	219583.0	33.49	0.345
Total lactation yield (kg)							
Parity 1	2050	0.136±0.04	58692.00	431420.51	372728.5	36.25	
Parity 2	1797	0.151±0.05	74758.78	493968.18	419209.4	37.41	
Parity 3	1533	0.158±0.06	74009.11	468391.50	394382.4	35.83	
Parity 4	1192	0.186±0.06	80433.88	432256.58	351822.7	35.61	
Parity 5	875	0.183±0.11	77444.34	423032.91	345588.6	35.05	
upto 5th parity	7446	0.109±0.03	51568.77	469023.25	292767.3	36.84	0.375
All parities	9003	0.103±0.02	47959.47	463693.05	296035.0	37.00	0.361
Lactation length (days)							
Parity 1	2050	0.061±0.04	432.07	6976.67	6544.6	29.33	
Parity 2	1797	0.074±0.04	489.66	6616.98	6127.3	28.74	
Parity 3	1533	0.065±0.05	383.73	5835.74	5452.0	27.11	
Parity 4	1192	0.184±0.07	1113.49	6025.87	4912.4	28.07	
Parity 5	875	0.118±0.08	636.01	5740.77	5104.7	27.58	
upto 5th parity	7446	0.071±0.02	471.44	6621.11	5288.8	28.93	0.201
All parities	9003	0.064±0.02	417.03	6515.57	5170.8	29.01	0.206
Dry period (day)	7283	0.033±0.01	551.01	16358.42	14886.7	49.55	0.090

estimated by four methods for first dry period were 0.14, 0.12, 0.09 and 0.09, respectively under univariate and multivariate REML analyses. The low magnitude of h^2 clearly indicated that this is influenced by the environmental factors and this trait may be improved by better feeding and management practices (Jakhar et al., 2017; Yadav and Singh, 2016).

The repeatability estimates for the dry period was 0.09 in the present study. Repeatability estimates for dry period in Pakistani buffaloes have been reported to range between 0.09-0.20 (Thevamanoharan et al., 2002; Salah-ud-Din, 1989). For Indian breeds estimates have also been higher (0.10 to 0.20) than the present study (Jamuna et al., 2015). Metry et al. (1994) reported dry period to have repeatability of 0.22 in Egyptian buffaloes while Ashmawy (1991) and Ayyat et al. (1996) reported estimates of 0.22 and 0.28, respectively.

Conclusions

Genetic parameters estimated for various productive traits in the present study were in the generally reported range of parameters for buffaloes. Milk yield was lowly heritable (10%) with high repeatability (35%). Genetic control of milk yield and lactation length was different for different parities. Genetic base needs to be widened to increase the genetic variation. Genetic selection of young bulls under artificial insemination program would be one segment requiring priority for better genetic gain.

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Authors' contribution

Research work was completed by MKB with the help of MSK and data collection and provision of literature by MIM and SUR and their valuable suggestion regarding design of experiment and data analysis.

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