GENETIC AND PHENOTYPIC TREND OF MILK PRODUCTION TRAITS AND AGE AT FIRST CALVING APPLYING RANDOM REGRESSION MODEL IN FRIESIAN CATTLE IN EGYPT

M.H. Mohamed¹, M.H. Khalil², M.M. Iraqi² and S.M. Zahed¹

1-Animal Production Research Institute, Agricultural Research Center, Ministry of Agriculture and Land Reclamation, Dokki, Giza, Egypt, 2- Department of Animal Production, Faculty of Agriculture at Moshtohor, Benha University, Egypt, Corresponding author: maher.khalil@fagr.bu.edu.eg

Received: 28/11/2019 Accepted: 26/3/2020

SUMMARY

The aim of this study was to detect genetic and phenotypic trends for test-day (TD) milk, fat and protein yields and age at first calving (AFC) in Frisian cattle in Egypt applying the random regression model (RRM). Data of 5237 TD milk yield traits were recorded for 953 Friesian cows, daughters of 208 sires and 944 dams from two herds in the Animal Production Institute (APRI), Egypt. Ten-month classes of lactation days were considered for the TD yields. The model included the random effects of direct additive genetic, permanent environment and error, while the fixed effects were herd-year-season of calving and parity, which was modeled by Orthogonal Legendre Polynomials. The additive genetic variance at first TD for milk, fat, protein yields and AFCwere4.7 kg, 13.3 g, 5.3 g and 2.8 monthrespectivelyand increased until the fourth, decreased until the lowest value at the tenth TD for milk but fat and protein yields the lowest value at the ninth TD. The lowest value of AFC was at the sixth TD then increased until the tenth TD. Heritability estimates at first TD were 0.12, 0.25, 0.25 and 0.05 for TDMY, TDFY, TDPY and AFC, respectively, and increased until the third for TDFY, TDPY and AFC but TDMY trait reached the highest value at the fourth TD, then decreased at the tenth TD.But reached the lowest value at sixth TD, and increased until the tenth TD. The phenotypic trendfor TDMY decreased by 26 to 18 kg from the first to the fifth TD, then it increased until the tenth TD. The phenotypic trend was started by 44 g and 18 g for fat and protein, respectively until the fourth TD then decreased until the ninth TD. The genetic trends were slightly positive for all traits, indicating that the selection program performs correctly.

Keywords: Test day milk yield, random regression model, genetic and phenotypic trends

INTRODUCTION

Random regression models (RRM) are currently used in the prediction of breeding values and in the estimation of the variance components for milk production traits of dairy cattle in several countries. Direct modeling of test day (TD) records instead of 305-d yields allows the shape of the lactation curve to be modeled with subsequently more precise adjustment for temporary environmental effects, avoidance of extended records for culled cows or lactations in progress, and evaluation of lactation persistency (Jamrozik and Schaeffer, 1997). In the basic structure of a RRM, the fixed part includes effects peculiar to all cows on the same test day and effects specific to cows on a given test day, such as pregnant or diseased, plus a factor accounting for the yield level on a specific day in milk (Ptak and Schaeffer, 1993) whereas individual lactation curves are fitted by random regression coefficients (Schaeffer and Dekkers, 1994; Jamrozik and Schaeffer, 1997). This feature of RRM allows for the prediction of breeding values and estimatingthe (co)variance functions throughout the whole lactation. Mean lactation curves are usually estimated on a large number of records and are characterized by

Issued by The Egyptian Society of Animal Production

quite regular patterns. As stated by Schaeffer 2004the use of either mathematical functions or fixed intervals of days in milk will generally lead to the same results. In this study, the trend is investigated at the phenotypic level by a fixed regression analysis of individual deviations around the mean curves for milk yield of first lactation Canadian Holsteins using some of the functions proposed to fit random effects in RRM for milk production traits in cattle. The aim of the present study was to estimate variance components, heritability and genetic and phenotypic trend for Friesian cattle raised in Egypt.

MATERIAL AND METHODS

Management and data structure:

A total of 5237 test-day milk, fat and protein yield records from 2000 to 2016 years were used in this study as shown in Table1.Cows were fed, on a ration consisting cotton seed cakes, barley wheat and rice bran, cows fed mainly on berseem and rice straw. In addition concentrates feed mixture from December to April. Mineral mixture bricks were offered *adlibitum* as soled minerals mixture in front animals, and on a balanced ration of a concentrates according to their production and weight. Limited amount of clover hay was used when available. Animals housed under open sheds covered with 3.5-4 meters high roofs. Heifers were bred for the first time when they reached 18-22 months or 350 kg body weight. Cows were artificially inseminated not before 60 days of calving using frozen semen from U.S.A and Canada. Cows milked two times a day by milking machine.

Data collected at monthly intervals in three experimental herds (Sakha and El karada in Kafrelshikh Governorate and El Serw in Damietta governorate) belonging to Animal Production Research Institute (APRI), Ministry of Agriculture, Egypt. Test-day milk yield (TDMY) records were measured based on an alternative am-pm monthly recording scheme. Milking was practiced twice a day at 7 am and 4 pm throughout the lactation period. In general, using TD models could have advantages over a 305-day model (Wilmink, 1987; Danell, 1982; Keown and VanVLeck, 1971). Cows with less than four TD records per lactation were excluded from the data set, while the maximum number of test day records per lactation was 10 records. Moreover, up normal phenotypic values of daily milk yield, fat and protein yield were removed from the dataset. The structure of the data analyzed is shown in Table1.

Table1. Structure of test day data analyzed for cattle's

Item	Data				
No. of sires	208				
No. of dams	944				
No. of base animals	953				
No of non-base animals	1560				
Total number of animals	2513				
Total number of lactation records	5237				

Measuring fat and protein percentages in milk:

Fat and protein percentages were measured by the method of infrared absorption automated spectrophotometry (Milk-o-Scan; Foss Electric, Hillerød, Denmark) at the Dairy Services Unit, Animal Production Research Institute, Sakha, Kafr El-Sheikh Governorate. The device needs a set of solutions: The first solution is used to wash the device after the analysis of the samples and before turning it off the name of this solution none foaming Stella 0.5% (Foss electric company, Denmark). The second solution is used to reset the device which gives the readings 0.000 so it is ready to read the new samples and its name is Triton x-100 and we use only 1 cm/liter of distilled water, and finally we have to give the device the order Prog. 2 then Prog.3 then Prog. 4, and then the device is programmed to read the cow milk samples. After that the percentages of fat and protein had been converted to yields in grams.

Statistical analyses:

Estimating variance (co) components using random regression model:

The monthly test-day milk yield (TDMY) were measured between 4 and 304 days in milk (DIM), divided into 10 classes. The first class included milk yield between 4 and 30 DIM, the second included milk yield between 31 and 60 DIM, and so on until the last class, which included milk yield between 270 and 304 DIM. The orthogonal polynomials of standardized units of time have been recommended as covariables(Kirkpatrick et al., 1990; Meyer, 1998). Orthogonal polynomials have computational advantages; the primary general advantage is the reduced correlations among the estimated coefficients.

The standardized unit of time, w, ranges from -1 to +1, was derived as:

$$t^{\bullet} = \frac{2(t - t_{min})}{t_{max} - t_{min}} - 1$$

Wheret_{min} is the earliest date (or the youngest age) and t_{max} is the latest date (or oldest age) represented in the data. The first six Legendre polynomial functions of standardized units of time are given in Table 2. Thus, if w = - 0.2, then the coverable that would go into the model (for order equal to 5) are shown in the last column of Table 2. Coverable based upon orthogonal polynomials are small numbers that reduce problems with rounding errors, and they provide relatively small correlations between the estimated regression coefficients.

Table2. The first six Legendre polynomial functions of standardized units of time

Order		For $w = -0.2$
0	$0.7071 w^0$	0.7071
1	$1.2247 w^1$	-0.2449
2	$-0.7906w^{0}+2.3717w^{2}$	-0.6957
3	$-2.8062w^{1}+4.6771w^{3}$	0.5238
4	$0.7955 w^{0}$ -7.9550 w^{2} +9.2808 w^{4}	0.4921
5	$4.3973w^{1}-20.5206w^{3}+18.4685w^{5}$	-0.7212

And so on, the first six Legendre polynomial functions can be put into a matrix of polynomial coefficients (Λ) as:

	0.7071	0	0	0	0	0)
	0	1.2247	0	0	0	0
$\Lambda' =$	-0.7906	0	2.3717	0	0	0
<u>n</u> –	0	-2.8062	0	4.6771	0	0
	0.7955	0	-7.9550	0	9.2808	0
	0	4.3973	0	-20.5206	0	18.44685)

Now we can define another matrix, M, as a matrix containing the polynomials of standardized time values. Legendre polynomials are defined within the range of values from -1 to +1. Thus, ages or time periods have to be standardized (converted) to the interval between -1 to +1. The formula is

$$t^{\star} = \frac{2(t - t_{min})}{t_{min} - t_{min}} - 1$$

c_{max} . this gives $\Phi = M \Lambda$. M=- ^tmin [] Orthogonal polynomials tend to reduce the correlations between estimated regression coefficients. This is advantageous, because the estimates would converge faster to the maximum or appropriate posterior distribution. There are other kinds of orthogonal polynomials, but Legendre polynomials are probably the easiest to calculate and utilize. Covariables based upon orthogonal polynomials are small numbers that reduce problems with rounding errors, and they provide relatively small correlations between the estimated regression coefficients.

Variance-covariance components were estimated by REML using the computer package VCE6 (Groeneveld*et al.*, 2010). The animal model was:

$$Y_{ijkl} = HYS_i + \sum_{m=1}^{4} \beta_{km}Z_{jlm} + \sum_{m=1}^{4} \alpha_{jm}Z_{jlm} + Pe_j + e_{ijkl}$$

where: Y_{ijkl} is the record *l*on trait within lactation made on herd-year-season (HYS) subclass i for the jth cow belonging to kth subclass (k ranged from 1 to 10 starting with k=1 and increased by 1 every 30 days thereafter along the trajectory from 4 to 304-d); HYS_i is the fixed effect of herd-year-season, Pe_j = random effect of permanent environment associated with all TD yields of the jth cow; β_{km} and α_{jm} = fixed and random regression coefficient, and e_{ijkl} = random residual effect associated with Y_{ijkl}.

The VCE6 program applying the Random Regression Model (RRM) was used to analyze the data using the Legendre polynomials method (Kirkpatrick *et al.*, 1990). The general RRM can be represented in matrix notation as:

Y = Xb + Za + Wp + e,

Where, Y = vector of observations on animal; b = vector of the fixed effects (HYS is the fixed effect of herd-year-season combination and P is the fixed effect of the lth parity (l = 12 levels for all parities); a = vector of solutions for additive genetic random coefficients; p = vector of solutions for permanent environmental random coefficients; e = vector of N different residuals; X, Z, and W = incidence matrices for fixed and random genetic and permanent environmental random effects, respectively. The

assumptions with respect to the components of the model were (Jamrozik and Schaeffer, 1997; Schaeffer, 2004):

$$\begin{pmatrix} a \\ p \\ e \end{pmatrix} \square N(0, V) \text{ where, } V = Var \begin{pmatrix} a \\ p \\ e \end{pmatrix} = \begin{pmatrix} G \otimes A & 0 & 0 \\ 0 & I\sigma_p^2 & 0 \\ 0 & 0 & R \end{pmatrix}$$

where G and P are (co)variance matrix of additive genetic and permanent environment random regression coefficients, respectively; A is an additive genetic relationship matrix among the cows; \bigotimes is a Kronecker product function; I is identity matrix and R is the diagonal matrix of temporary environmental variances. The mixed model equations for this model would be:

$$V = \begin{pmatrix} X'X & X'Z & X'W \\ Z'X & Z'Z + G^{-1} \otimes A^{-1} & Z'W \\ W'X & W'Z & W'W + I \otimes P^{-1} \end{pmatrix} \begin{pmatrix} b \\ a \\ p \\ p \end{pmatrix} = \begin{pmatrix} X'Y \\ Z'Y \\ W'Y \end{pmatrix}$$
$$E \begin{bmatrix} y \\ a \\ p \\ e \end{bmatrix} = \begin{bmatrix} Xb \\ 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}; \text{ and } V(p) = K_p \otimes I;$$
$$V(e) = R,$$

Where k_a and k_p are the genetic and permanent environmental covariance matrices between random regression coefficients, respectively. A is the additive genetic relationship matrix; I is an identity matrix, and R represents a diagonal matrix containing the residual variances.

The genetic (G) and permanent environmental (P) covariance between test-days were estimated using:

$$G = (1 \ ti \ ti^2 \ \dots) Ka \begin{bmatrix} 1 \\ tj \\ tj^2 \\ \dots \end{bmatrix} and P = (1 \ ti \ ti^2 \ \dots) Kp \begin{bmatrix} 1 \\ tj \\ tj^2 \\ \dots \end{bmatrix}$$

Heritabilities (h^2) are computed using the package of VCE6 as (Groeneveld *et al.*, 2010):

$$h^2 = \frac{\sigma_{gi}^2}{\sigma_{gi}^2 + \sigma_{pei}^2 + \sigma_{ei}^2}$$

Where: σ_{gi}^2 is the additive genetic variance of the ith TD; σ_{pi}^2 is the permanent environmental variance and σ_{ei}^2 is the residual variance.

Predicting the breeding values of the genetic trend:

Predicted breeding values (PBVs) for cows were estimated using the computer package of PEST program (Groeneveld *et al.*, 2001) for test day milk, fat and protein yields and age at first calving according to the following model:

$$y = Xb + Za a + Zc c + e$$

where: y = Vector of observations, X = Incidence matrix relating fixed effects to y, b = Vector of an overall mean and fixed effects, (HYS is the fixed effect of herd-year-season combination and P is the fixed effect of the 1th parity (l = 12 levels for all parities), Za = Incidence matrix relating direct additive genetic effects to y, a = Vector of random effect (direct additive genetic associated with the

incidence matrix Za), Zc = Incidence matrix for permanent environmental effect, c = Vector of permanent environmental effect associated with the incidence matrix Zc and e = Vector of random residual effects N (0, $I\sigma^2_{e}$); I is an identity matrix.

Solutions for the equations of animals were computed from the pedigree file, one animal at a time for animals with records and animals without records (sires and dams). A diagonal element (d_t) and an adjusted right-hand side (V^*_t) were accumulated with each pedigree file record for the tth animal. For the animals with and without records, the formula used to estimate the PBV was that of (Kennedy 1989):

$PBV = [_{t}^{y}/d_{t}]$

The predicted error variances (PEV) of predicted breeding values (PBVp) were estimated for each individual as: $PEV_p = d_j \sigma^2_{e}$ (Korsgaard *et al.*, 2002); where d_j is the jth diagonal element of inverse of the appropriate block coefficient matrix and σ^2_{e} is the residual variance. The accuracy of PBV for each individual was estimated according to Henderson (1975) as:

$$r_{A} = \sqrt{1 + F_{j} - d_{j}\alpha_{a}}$$

Where

The accuracy of prediction of the ith animal's breeding value; F_j=inbreeding coefficient of animals (assumed equal to be zero); d_j was defined before; and $\alpha_a = \sigma_e^2 / \sigma_a^2$.

Plotting the genetic and phenotypic trends:

The phenotypic trend was measured as the regression of least squares means on years. The breeding values of the animals with records and without records were estimated using the PEST program (Groeneveld*et al.*, 2001). Accordingly, the genetic trend was measured by regressing the breeding values on years.

RESULTSANDDISSCUTION

Means and variations:

Number of observations, means and their standard deviations and coefficients of variation (CV) for thetraits studied are shown in Table 3.

Table3. Number of observations, means, and standard deviations (SD) and coefficients of variation (CV%) for test day (TD) milk, fat and protein yields and age at first calving (AFC)

	No. of													
TD	obs. for	Т	DMY		TDFY			Т	TDPY			AFC		
NO.	all milk	Mean	SD	Cv	Mean	SD	Cv	Mean	SD	Cv	N. of	Mean	SD	Cv
	traits	kg	kg	%	Kg	kg	%	kg	Kg	%	obs.	Mo.	Mo.	%
1	809	10.4	5.6	46	0.43	0.2	55	0.32	0.2	51	533	32.2	5.8	15
2	861	11.3	6.6	45	0.43	0.2	50	0.31	0.2	51	566	32.3	5.7	15
3	870	12.9	5.2	44	0.45	0.3	56	0.31	0.2	50	573	32.2	5.8	15
4	866	11.4	5.1	45	0.39	0.2	50	0.29	0.1	47	566	32.2	5.8	15
5	621	11.1	5.8	43	0.40	0.2	51	0.28	0.1	46	384	32.2	4.3	13
6	457	10.4	5.6	44	0.35	0.2	53	0.27	0.1	48	268	32.4	4.4	14
7	331	10.2	5.6	45	0.36	0.2	55	0.26	0.1	48	194	32.2	5.5	14
8	227	10.6	5.6	48	0.33	0.2	54	0.24	0.1	48	123	32.2	5.6	14
9	135	9.2	5.6	50	0.33	0.2	53	0.23	0.1	48	73	32.2	5.8	15
10	59	9.4	5.9	53	0.32	0.2	58	0.25	0.1	57	32	32.3	5.7	15
TD-	TD - Test day, TDMV - Test day milk yield TDEV - Test day for yield TDEV - Test day protein yield and AEC - A ge at													

TD = Test-day, TDMY = Test-day milk yield, TDFY = Test-day fat yield, TDPY = Test-day protein yield and AFC = Age at first calving.

Means and standard deviations of test-day milk yield increased from the first test day (TDMY1) to the peak in the third test day (TDMY3) which was 12.9±5.2 kg. TDM decreased from the fourth and fifth TDMY which were11.1±5.8 kg and fixed from the sixth TDMY to the eighth test day which were 10.6±5.6 kg and in the lasttwotest daywere 9.4±5.9 kg (Table3). Means and standard deviations of test day fat yield were fixed from the first (TDFY1) to the seventh test day fat (TDFY7) which was 0.36 ± 0.2 g. TDFY decreased from the eighth TDFY8 to the last TDFY10 were $0.32g\pm0.2g$.(Table3).For which protein-test day yield, the means were fixed from the first(TDPY1) to the fifth test day protein (TDPY5) which was 0.28±0.1 g. TDPY decreased from the sixth TDPY6 to the last TDPY10 which were 0.25 ±0.1 g. For age at first calving (AFC) the mean was nearly constant to be 32±5 month. Coefficientsof variation ranged from 43% to 53% for TDMY, from 50% to 58% for TDFY, from 46% to 57% for TDPY and from 0.13 to 0.15 for age at first calving.

Variances:

Estimates of additive genetic(V_A), permanent environmental(V_{Pe}), phenotypic (V_P)and résiduel (V_E)variances are presented in Figures (1), (2), (3) and (4) for test-day milk, fat, protein yields and age at first calving(AFC). The additive genetic variance (V_A) estimates at first test day were 4.7 kg, 13.3 g, 5.3 g and 2.8 month then they increased at the fourth test day to be 7.6 kg, 38.7 g, 14.3 g and 3.5 month and decreased thereafter, reaching the lowest value at the tenth test day for milk yield (3.4 kg) and at the tenth test day for fat, protein yield and age at first calving 13 g, 7.7 g and 5.7 month. Similar results have been reported by (Biassus et al. 2011) and (El Faro 2012)reported that the genetic variance values ranged from 2.3 to 5kg, from 1 to 7.5g and from 1 to 5g for TDMY, TDFY and TDPY, respectively.

Permanent variance estimates(**Vpe**) rangedfrom4.6 to 12.7 kg, 30.9 to 69g, 10.6 to 23.9 g and 2.6 mo to 8.6 mo, for TDMY, TDFY, TDPY and AFC respectiviley. Simler trends werealsoreported by (Jamrozik and Schaeffer 1997) and (Jensen *et al.* 2001) in cattle.The residual variance estimâtes(Ve) for all traits rangedfrom 8kg to 11kg, 0 to 8g, 1g to 4g and from 16 to 23mo for milk, fat, protien and AFC traits, respectively.

The phenotypic variance (V_P) estimates for milk, fat, protein and age at first calving traits rangedfrom 19.2 to 26.3 kg, 45.6 to 113g, 17.9 g to 41.5g and 24.7 mo to 34.8 mo, respictively. The phenotypic variance for the milk testdaydecreased until TD6 which it was 19.6 kg then increased until the TD10 which it was 23.9 kg. The

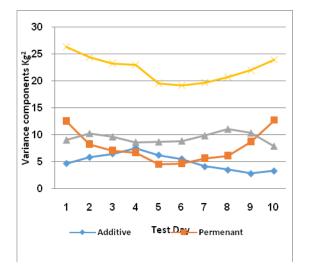


Fig. 1.Estimates of additive genetic (v_A) , permanent environmental (V_{Pe}) , residual variances (V_E) and phenotypic variances (V_P) for test day milk yield (kg).

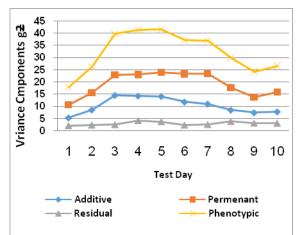


Fig. 3. Estimates of additive genetic (V_A) , permanent environmental (V_{Pe}) , residual variances (V_E) and phenotypic variances (V_P) for test day protein yield (g).

Heritability values:

Heritability estimates for TDMY at selected TD are shown graphically in (Figure 5). Estimates were

fat test-dayincreaseduntil TD4, itwas 113.1 g thenitdecreaseduntil TD7, itwas99,9gthendecreaseduntil TD10, itwas 55.2 g. Thesame trend wasobtained for protien trait. For AFC trait V_Adecreaseduntil TD6, 21.27 mo thenitincreaseduntil TD7 to be 24.12 mo, thenincreaseduntil TD10 whichreached 24.7 mo. Theseresultswere agreementwiththose of (Biassus et al. 2011)whoshowedthat the values of phenotypic variances rangedfrom 20 to 27 kg, 25 to 35 g and 20 to 35 g for TDMY, TDFY and TDPY respictively. (El Faro et al 2012)revealedthat the phenotypic, permanent and genetic variances rangedfrom 2 to 7 1.8 to 5 and 0.2 to kg, 1.2 kg for TDMY.Theseresultswerenearlysimilar to thosereportedby(Cobuci et al 2005) permanent and geneticvariaces values rangedfrom 19 to 29 kg and 8 to 20 kg for TDMY respectiviley.

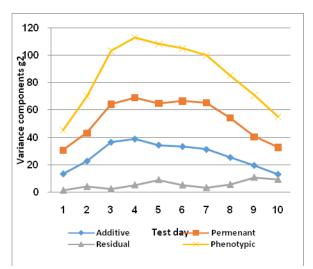


Fig. 2. Estimates of additive genetic (v_A), permanent environmental (V_{Pe}), residual variances (V_E) and phenotypic variances (V_P) for test day fat yield (g).

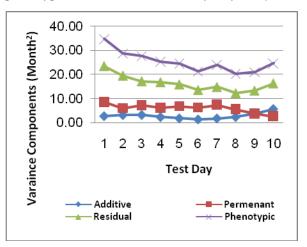


Fig. 4. Estimates of additive genetic (V_A) , permanent environmental (V_{Pe}) , residual variances (V_E) and phenotypic variances (V_P) for age at first calving (AFC).

low at the beginning of the test day (0.17), and gradually increased reaching the highest value at the fourth test day (0.33) and decreased gradually until

they reached the lowest value at the tenth test day of lactation (0.14). The heritability estimates for fat and protein yields showed the same trend where the estimates were 0.29 and 0.3 on the first test day and reached 0.35 and 0.36 at the fourth test day, and finally decreased at the ninth test day in fat (0.24) and at the tenth test day in protein yield (0.29). Similar trends were reported by Rosati and (Van Vleck,2002)for milking cows. Heritability estimates for AFC were low at the beginning of the test day (0.07), and gradually increased, reaching the highest value at the third test day (0.12) and decreased in the sixth test day (0.07), and then it increased until the tenth test day whenit was 0.24. In general, heritability estimates for traits had wide ranges and tended to increase toward the edges of the defined lactation trajectory. Most heritability estimates obtained by RRM were high at the edges as stated by(Jamrozik and Schaeffer, 1997)in dairy cows. Difficulties in the model in getting acceptable variances at the extremes of the lactation can be explained, in part, by the biological processes that occur at the beginning of lactation and the smaller number of records at the end. Jamrozik and Schaeffer (1997) and El- Saied (2004) pointed out that these parametric functions tend to overestimate the genetic variances and underestimate the genetic correlations among milk yield at the beginning and the end of lactations.

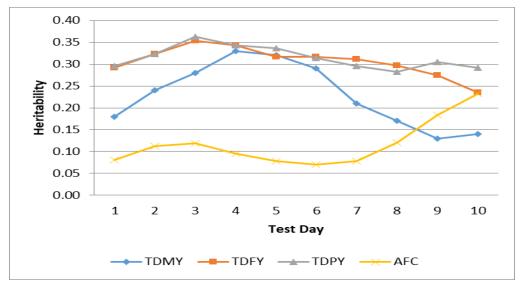


Fig. 5.Estimates of heritability of test-day milk (TDMY), fat (TDFY) and protein (TDPY) yields and age at first calving (AFC).

Predicted breeding value (PBV):

The minimum and maximum predicted breeding values (PBV) and their accuracies for milk, fat and protein yields and age at first calving are given in Table 4. The PBV for milk, fat and protein yields and age at first calving ranged from -1297 to 1896kg, -5.9 to 6.2 kg and -7.1 to 7.5kg and -2.6 to 2.4 month, respectively. Using TD animal model methodology in cattle, (Zutere, 2008) found that the estimated

breeding values for milk, fat and protein ranged from -1014 to 1966kg, from -40.75 to 93.59kg and from -37.33 to 59.86kg, respectively. The accuracies ($^{rA^{}}$) of minimum and maximum estimates of PBV were high in all traits (Table 4). This may be due to that estimate of heritability were highly associated with more available pedigree information for all individuals (Korhonen, 1996; Korsgaard *et al.*, 2002).

Table 4.Minimum, maximum and ranges of predicted breeding values (PBV), predicted error variance (PEV) and accuracy of prediction($^{rA^{\wedge}}$)for test day milk yield (TDMY), fat and protein yields in Friesian cattle raise in Egypt

		Minimum			Range in		
Trait	PBV PEV ^{A^}		PBV PEV		rA^	PBV	
TDMY, kg	-1296.5	324.9	0.99	1895.6	634.0	1.00	3192.1
Fat yield, kg	-5.9	4.1	0.89	6.2	5.2	0.92	12.1
Protein yield, kg	-7.1	4.6	0.91	7.5	5.8	0.93	14.6
Age at first calving, mo.	-2.6	1.2	0.65	2.4	1.7	0.78	5.0

Phenotypic trend:

Phenotypic trend for TDMY, TDFY and TDPY and age at first calving are shown in Figure 6. Phenotypic trends for TDMY, TDFY, TDPY and AFC were decreased from TD1 to TD10. The decrease in phenotypic trend in all traits may be attributed to low nutritional level applied and management practices in different herds.

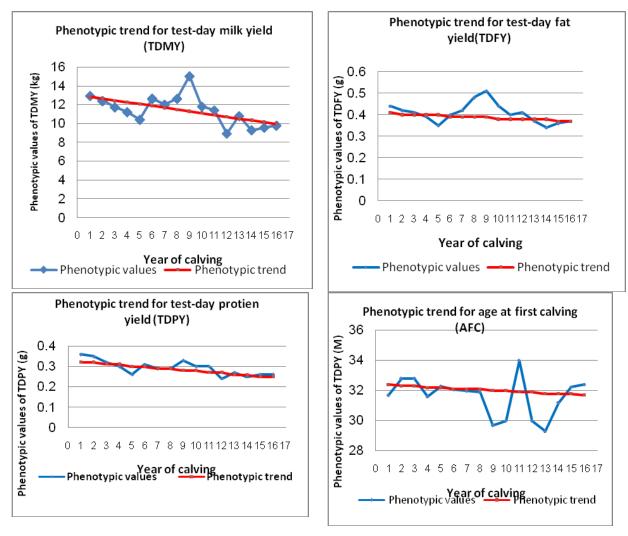
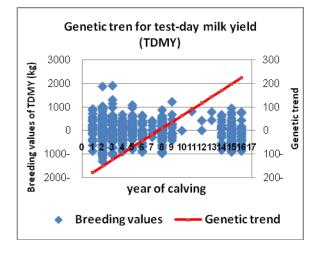


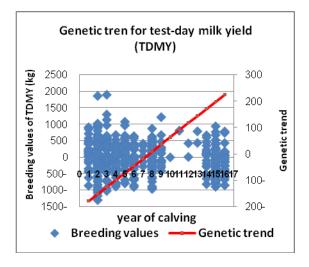
Fig. 6.Phenotypic trend for test-day milk, fat, protein yields and age at first calving in Friesian cattle raised in Egypt.

Genetic trend:

Genetic trend for TDMY, TDFY, TDPY and age at first calving are shown in Figure7, the values of genetic trend for TDMY, TDY, TDPY, and AFC traits increased from TD1 to TD10. These results indicated that improvement program of selection was



practiced in the farms. Similar trends were also reported by some investigators e.g.(Yaeghoobi *et al*2011) and (Katok and Yanar 2012),(Muller and Botha, 2003) for TDMY, (Jamrozik and Schaeffer, 1997).



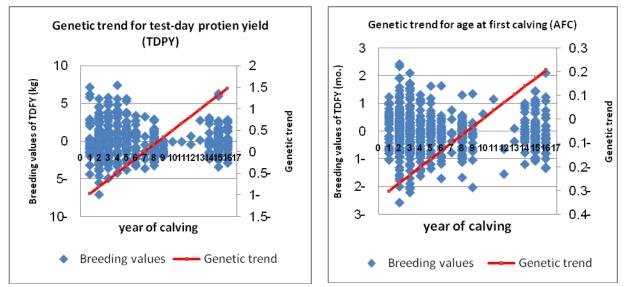


Fig. 7. Genetic trend for test-day milk, fat, protein yields and age at first calving in Friesian cattle raise in Egypt

CONCLUSIONS

- 1) The test-day milk yield during the first three to five months of lactation could be adopted as an early selection criterion to increase milk yield.
- 2) Random regression model (RRM) was considered to be efficient model in detecting the fluctuations in genetic variance during the lactation period. It would permit better modeling for repeated records throughout the lactation period and could be chosen as an accurate method for predicting breeding values.
- **3)** Genetic trend showing the improvement of testday milk, fat, protein yields and age at first calving.So selection program was practiced correctly in this herd.

4)For all traits, the phenotypic trends showing deteriorating trends indicating the presence of some environmental inadequacies especially for nutritional level.

REFERENCES

- Biassus I.O, JA. Cobuci, C.N. Costa, P.R.N. Rorato, J.B. Neto, and L.L. Cardoso, 2011.Genetic parameters for production traits in primiparous Holstein cows estimated by random regression models. R. Bras. Zootec., v.40, n.1, p.85-94.
- Danell, B. 1982. Studies on lactation yield and individual test-day yield of Swedish dairy cows: III Persistency of milk yield and its correlation with lactation yield. Acta. Agriculture Scandinavica, 32:93-101.
- El Faro L, Cardoso V. L and de Albuquerque L. G 2012.Variance component estimates applying random regression modelsfor test-day milk yield in Caracu heifers. Genetics and Molecular Biology, 31, 3, 665-673
- El-Saied, U. M., 2004. Random regression estimation of genetic parameters for milk yield and protein test day model for percentage of

primiparous Frisian cows in Egypt. Egyptian J. Anim. Prod., 41 (1): 1-10.

- Groeneveld, E., KovačM, and MielenzN, 2010.VCE 6, Users guide and reference manual, Version 6.0.2.
- Groeneveld, E., Kovac M., andWang T, 2001. PEST, Users guide and reference manual, Version 4.2.3.
- Henderson, C.R. 1975.Best linear unbiased estimation and prediction under a selection model. Biometrics 31(2): 423-447.
- Jamrozik, J., Schaeffer, L.R., 1997.Estimates of genetic parameters for a test day model with random regressions for yield traits of first lactation Holsteins. J. Dairy Sci. 80:762-770.
- Jensen, J., Jamrozik J. and Schaeffer L. R, 2001.Modelling production in all lactations in dairy cattle using random regression test day models. Book of Abstract proc. 52nd Annu. Mtg.of the EAAP. August, 26-29,2001,Budapest.
- Katok, N. and Yanar M 2012. Milk traits and estimation of genetic, phenotypic and environmental trends for milk and milk fat yields in Holstein Friesian cows. Int. J. Agric. Biol., 14: 311–314.
- Kennedy, B. W. 1989. ANIMAL MODEL BLUP -Crasmus intensive graduate course. University of Guelph, Dublin.
- Keown, J. F. and VanVleck, L. D. 1971.Selection on test-day fat percentage and milk production. J. Dairy Sci., 54:199-203.
- Kirkpatrick, Lofsvold M., D., and Bulmer M., 1990.Analysis of the inheritance, selection and evolution of growth trajectories.Genetics., 124: 979-993.
- Korhonen, T. (1996)."The dairy cattle evaluation of 1996.http://www.mloy.fi/faba/blup/blup1.html,23 .01.1977(h:14:13)(Article).
- Korsgaard, I.R.; Andersen, A.H. and Jensen, J. 2002. Prediction error variance and expected response to selection, when selection is based on the best predictor for Gaussian and threshold characters,

traits following a Posiion mixed model and survival traits. Genet. Sel. Evol., 34:307-333.

- Meyer, K. 1998b. Modeling repeated record: Covariance function and random regression models to analyze animal breeding data. 6th WCGALP. 25:512-520. Armidale, Australia.
- Muller, C.J.C., and Botha J.A, 2003. The response to selection during first lactation on the phenotypic and genetic trends in the Elsenburg Holstein-Friesian herd. South African Journal of Animal Science, 33 (2): 111-116.
- Ptak, E., Schaeffer, L.R., (1993).Use of test day yields for genetic evaluations of dairy sires and cows.Livest. Prod. Sci. 34:23-34.
- Rosati, A., and L. D. Van Vleck, 2002.Estimation of genetic parameters for milk, fat, protein and Mozzarella cheese production in Italian river buffalo population. Livest. Prod. Sci. 74:185–190.
- Schaeffer, L.R., 2004.Applications of Random Regression models in animal breeding.Livest. Prod. Sci. 86:35-45.

- Schaeffer, L.R., Dekkers, J.C.M., 1994. Random regressions in animal models for test day production in dairy cattle. Proc 5th World Congr.Genet.Appl. Livest. Prod., Guelph, ON, Canada, 18:443-446.
- Takma c. and Akbas y, 2007.Estimates of genetic parameters for test day milk yields of a Holstein Friesian herd in Turkey with random regression models. Arch. Tierz., Dummerstorf50 4, 327-336.
- Wilmink, J. B. M. 1987. Efficiency of selection for different cumulative milk, fat and protein yields in first lactation. J. Dairy Sci., 17:211-224.
- Yaeghoobi, R., Doosti A, Noorian A.M and Bahrami A.M, 2011.Genetic parameters and trends of milk and fat yields in Holstein's dairy cattle of west provinces of Iran. Int. J. Dairy Sci., 6: 142–149.
- Zutere, R. 2008.Estimates of breeding values for dairy cattle using test-day milk yields.Agronomijas Vestis (Latvian Journal of Agronomy), 10: 293-299.

الإتجاه الوراثى والمظهرى لصفات إنتاج اللبن والعمر عند أول ولادة بتطبيق نموذج الإنحدار العشوائى فى ماشية الفريزيان فى مصر

محمد حماده محمد'، ماهر حسب النبي خليل'، محمود مغربي عراقي'، سميح محمد زاهد'

ا - معهد بحوث الإنتاج الحيواني، مركز البحوث الزراعية، ٢ - قسم الإنتاج الحيواني، كلية الزراعة بمشتهر، جامعة بنها

الهدف الرئيسى من هذه الدراسة هو تحديد الإتجاهات الوراثية والمظهرية لصفات إنتاج اللبن ومكوناته من دهن وبروتين وصفة العمر عند اول ولادة باستخدام بيانات يوم الإختبار وذالك بتطبيق نموذج الإنحدار العشوائي وذالك لماشنة الفريزيان المرباة في مصر إشتملت الدراسة على ٢٣٧ سجل من ٩٥٣ بقرة التي هي بنات لـ ٢٠٨ طلوقة و ٩٤٤ أم من قطيعين للأبقار الفريزيان التابعين لمعهد بحوث الإنتاج الحيواني بمصر تم تقسيم موسم الإنتاج إلى ١٠ أجزاء حيث من كل شهر نسجل يوم حليب (يوم الإختبار) شمل نموذج تحليل البيانات بعض التأثيرات العشوانية وهي التأثير المضيف للأبقار والطلائق وكذالك التأثير البيئي الدائم وشملت التأثيرات الثابتة فصل وسنة الولادة وكذلك القطيع معا ثم موسم الولادة وكانت قيم التمينيف للأبقار والطلائق وكذالك التأثير البيئي الدائم وشملت التأثيرات الثابتة فصل وسنة الولادة وكذلك القطيع معا ثم موسم الولادة وكانت قيم التباين الوراثي المضيف في أول يوم إختبار لصفة إنتاج اللبن والدهن والبروتين وكذالك العمر عند أول ولادة كالاتي على التوالى ٢,٤كم و وحرم و ٢,٥مم و ٢,٢ شهر وزادت حتي يوم الإختبار لمانية والدهن والبروتين وكذالك العمر عند أول ولادة كالاتي على الإختبار أقل لصفة أنتاج الدهن والبروتين حيث وصلت أقل قيمه في الإختبار التاسع ولكن لصفة العمر عند أول ولادة ولقل قيمة لها عند يوم الإختبار وللعد ثم أخذت في الزيادة حتي يوم الإختبار العاشر. وكانت قيم المكافئات الوراثية لصفات انتاج اللبن والدهن والبروتين والعمر عند أول ولادة ولي لصفة أنتاج الدهن والبروتين حيث وصلت أقل قيمه في الإختبار التاسع ولكن لصفة العمر عند أول ولادة ولقل قيمة لها عند يوم الإختبار والسدس ثم أخذت في الزيادة حتي يوم الإختبار العاشر. وكانت قيم المكافئات الوراثية لصفات انتاج اللبن والدهن والبروتين والعمر عد أول ولادة ول يوم إختبار كانت ١٢، و٢٠، و٢٠، و٢٠، على العاشر. وكانت قيم المكافئات الوراثية لصفات انتاج اللبن لوالا لحفات إلى في والده و٢٠ ولاد ول ول يوم إختبار الدائل لحفات إلغان والدهن أعلى قيمة عند يوم الإختبار الرابع ثم بعد ذالك أخذت جميع الصاف في في أول يوم إختبار الحاشر، وتراوحت قيم التوالي وبدأت بعد ذالك في التزايد حتي يوم الإختبار الثالث لصفات إلى والدهن ول يوم عنه أول ولادة ولكن صفة إلغربار العاشر. و٢٠، على التوالي في أعلى قيم عند يوم الإختبار الأول حتى يوم الإختبار والعمر