

GENETIC EVALUATION OF THREE QUAIL LINES\ BY FULL DIALLEL CROSS DESIGN II- EGG PRODUCTION

JAMEELA H. SALIH and SHEKHMUOU H. HUSSEN

Dept. of Animal Production, College of Agriculture, University of Duhok, Kurdistan Region - Iraq

(Received: April 16, 2018; Accepted for Publication: December 2, 2018)

ABSTRACT

A total of 1307 quail chicks were hatched in two lots (450 and 857 chicks), from three quail lines (White-W, Light brown-Lb and Dark brown-Db). The both sexes of each line mated as full diallel cross design, and 9 genotypes (WW, WLb, WDb, LbW, LbLb, LbDb, DbW, DbLb, DbDb) were resulted. The unsexed chicks from each genotype were reared during the growth period from (1-35 days old) in cages (with dimensions of 85X85X85 cm.) with two replicates. Then at 35 days old, the birds were sexed and redivided on the same cages with two replicate also, for each genotype separately as families (15 females and 5 males). The experiment was conducted at the Poultry Farm of Animal Production Dept., College of Agriculture, University of Duhok, Kurdistan region-Iraq. The trial aimed to evaluate the genetic background of three quail lines for egg production traits by using the full diallel cross design. The following genetic parameters were estimated; general combining ability (GCA), specific combining ability (SCA), heterosis (H%), reciprocal effect (R_E), maternal effect (M_E), genetic value (GV), heritability (h^2) and genetic correlation (r_g); on egg production (EP) traits (egg weight EW, egg number EN and egg mass EM). The main results from the present study could be summarized as follow: SCA resulted as significant differences for all EP traits. All genotypes didn't significantly differ in heterosis percentage for egg weight and egg number. Reciprocal effect resulted in a significant ($p \leq 0.05$) effect for all EP traits. As conclusion, it may use the Db line as dam for egg production traits.

KEY WORDS: Quail, diallel cross, genetic parameters, layer performance.

INTRODUCTION

The quail birds are valuable resources for meat and eggs due to their early sexual maturity age, high annual egg numbers (250–270 eggs), resistance to common poultry diseases and hard conditions, in addition to more persistence for egg production with high level up to 1.5 years (Murakami and Ariki, 1998). However, Piao et al. (2004), reported that the eggs number in quail was insignificantly differed among different genetic groups at different periods. The full diallel cross design often utilizes the same parents as females and males which make the design a little complicated (Isik, 2009). Mahipal, et al. 2001, used diallel cross in three quail lines and showed that the variance due to GCA was more important for egg production and egg weight, but variance due to SCA was significant for age at first egg and reproductive traits. Analysis of combining ability

for (4) quail lines (L1, L2, L3 and L4) using diallel crosses was conducted by (Drumond, et al. 2014 and 2015), they represented significant effect for GCA on weekly egg number and egg yield, while the SCA was insignificantly affecting weekly egg number, egg yield and egg weight characteristic. Amin, et al. (2013) estimated maternal effect in crossed of 2x2 diallel mating and used two local strains (Egg line and Meat line) and two crosses, they found that the maternal effect estimates had highly significant values and egg line was better as sire than meat line. Sezer, (2007) estimated the genetic and phenotypic parameters for some characteristics of Japanese quail and observed that the heritability of egg weight was high (0.83), he added that the heritability of all other characteristics of egg quality which related to the egg weight (size) trait, were high too. However, the heritability estimate of egg weight in Japanese quail was ranged from

0.18 to 0.68 (Koack, et al. 1995; Saatci, et al. 2006 and Khaldari, et al. 2010; Anang, et al. 2000; Dana, et al. 2011 and Hussen, et al. 2016). Moreover, heritability estimate of egg number was ranged from (0.09 to 0.48) in layers (Luo, et al. 2007 and Dana, et al. 2011). Hidalgo et al. (2011) suggested that in case of positive genetic correlation coefficients between EW and EN traits, the selection for one characteristic will improve another one, but in case of negative genetic correlation coefficients both traits should be involved in selection program.

(Lotfi, et al. 2012) estimated the genetic correlation between egg production characteristics and showed that the quail had high positive genetic correlation coefficients between egg weight at various weeks of age (0.88-0.99) and between egg number at different periods (0.98-0.99). This research aimed to investigate the genetic parameters of egg production traits using full diallel cross design.

MATERIAL AND METHODS

Place and design of Experiment:

A total of 1307 quail birds from two lots (450 and 857 chicks), were used in this research which included, White (W), Light brown (Lb) and Dark brown (Db) lines. The trial was conducted at the Poultry Farm of Animal Production Department, College of Agriculture, University of Duhok, Kurdistan region-Iraq.

The unsexed chicks from each genotype of the progeny were reared during the growth period from (1-35 days old) in cages (dimensions of 85X85X85 cm.) with two replicates. Then at 35 days old, the birds were sexed and redivided on the same cages with two replicate also, for each genotype separately as families (15 females and 5 males).

Environment and feeding

The progeny flock birds were hatched in two lots. The average temperature inside the house was (35 °C) in the first week, then decreased gradually by (2 °C) per week, up to 8 weeks old. The average relative humidity inside the rearing house was (31 %) during the whole study period (ranged from 21 % - 39 %). Light program included 23 hours/day for the first week of age, then modified to be 15 hours/day from the second week old until the end of trial (from 5 am to 8 pm lightness

and from 8 pm to 5 am, darkness) Feed was offered *ad libitum* manually for both generations (lines and crosses), which included three rations; starter (2850 K. cal. ME/ kg & 26 % CP) from (0-4) weeks old, grower (2850 K. cal. ME/ kg & 21% CP) from (4-6) weeks old and layer or breeder (2737 K. cal. ME/kg & 15.7% CP) from 6 weeks old-end of the trial, respectively, according to (Lesson and Summers, 2005).

STUDIED PARAMETERS:

The following genetic parameters were studied for EP traits; and were estimated according to (Falconer, 1988 and Williams, et al. 2002) in the crosses (progeny) with avoiding sire effect:

General Combining Ability (GCA): The values of GCA for the lines were computed as the means of specific line for giving trait.

$$GCA_i = \Sigma y_i / n$$

Where: GCA_i= the GCA for line _i; y_i = trait for a progeny from the specific line _i. n= the number of all progenies.

Specific Combining Ability (SCA): The values of SCA for the crosses were computed as the difference between average of cross with its reciprocal cross and average of GCA for both lines, for given trait according to the following formula:

$$SCA_{AB} = [(AB + BA) / 2 - (GCA_A + GCA_B) / 2]$$

Where: AB= the cross; BA= reciprocal cross.

Heterosis (H %): Hybrid vigor (heterosis) was computed on the basis of percentage of mid-parents for the given trait as following equation:

$$H \% = \{F1 - [(P1 + P2) / 2] / [(P1 + P2) / 2] \times 100\}$$

Where F1 = mean of the first generation and (P1 and P2) are the parents in diallel cross design.

Reciprocal Effect (R_E): Reciprocal effect was computed as half of the difference between the cross and its reciprocal cross for the given trait, according to the following formula:

$$R_E = (y_{ji} - y_{ij}) / 2$$

Where: y_{ji} = reciprocal cross; y_{ij} = the cross

Maternal Effect (M_E): Maternal effect was computed as the mean deviation of progeny for a particular dam, from mean estimated for a particular sire, according to the following formula:

$$M_E = (\bar{y}_{.i} - \bar{y}_i)$$

Where: $\bar{y}_{.i}$ = particular dam average; \bar{y}_i = particular sire average.

Genetic Value (GV): Genetic value of a cross is representing the GCA of both parents (lines) and SCA

of the same cross, and is calculated according to the following formula:

$$GV_{(AB)} = GCA_{(A)} + GCA_{(B)} + SCA_{(AB)}$$

Heritability (h^2): Heritability estimates were computed from the relationship between the average of the one-parent and the offspring (from regression coefficient), and from the variance components which were resulted from the effect of both GCA and SCA as random factors in ANOVA. So, it could summarize the computation of heritability as the following two equations:

$$h^2_1 = 2b; \text{ And } h^2_2 = 4(\sigma^2_{GCA} + \sigma^2_{SCA}) / \sigma^2_P$$

$$h^2 = (h^2_1 + h^2_2) / 2$$

Where b= regression coefficient of offspring on parents; σ^2_P is a phenotypic variance of a trait, and it included additive, dominant and residual effects.

Genetic Correlation (r_g): Genetic correlations between some previous traits were calculated according to the geometric equation as follows:

$$r_g = \sqrt{(CovZ2X1 * Cov Z1X2) / \sqrt{(CovZ1X1 * Cov Z2X2)}}$$

Where: Z= the observations on parent; X= the observations on offspring; 1= the first trait; 2= the second trait.

Statistical analysis:

The experiment were designed as diallel cross within completely randomized design (CRD), and collected data was analysed using SAS (SAS

Institute, 2010) software via mixed model for the effects of GCA and SCA as random factors in addition to the fixed effects of the replication (lot) was applied to analyse the previous studied traits in order to estimate the heritability according to the following model,

$$Y_{ij} = \mu + R_i + GCA + SCA + e_{ij}$$

Where: Y_{ijkl} : the observations of the studied trait; μ = overall mean; R_i : The effect of replication; GCA: the random effect of GCA; SCA: the random effect of SCA, e_{ij} : random error; i: lot (Replication); j: individual (bird). The differences between the means were analysed using Duncan multiple range test (Duncan,1955).

RESULTS AND DISCUSSION

General Combining Ability (GCA):

Table (1) illustrating the GCA values of egg number, egg weight and egg mass. The results showed that there are no significant differences among studied lines. However, pure W line gave the highest insignificant values of GCA at first and second weeks of laying for egg number (4.53 & 5.07 eggs), egg weight (10.10 & 10.46 gm) and egg mass (45.77 & 53.00 gm) respectively, compared with both brown (Lb and Db) lines.

Table (1). General combining ability for egg characteristic at first two weeks of laying period.

Lines	Egg number/hen.		Egg weight (gm)		Egg mass (gm)	
	W1	W2	W1	W2	W1	W2
W	4.53±0.29	5.07±0.17	10.10±0.01	10.46±0.46	45.77±2.99	53.00±2.00
Lb	4.25±0.34	4.56±0.24	9.92±0.08	10.36±0.13	42.14±3.69	47.37±3.15
Db	4.21±0.49	5.02±0.3	10.02±0.06	10.48±0.04	42.19±4.56	52.69±3.34
Sig.	Ns	Ns	Ns	Ns	Ns	Ns

Ns = Non-significant; values = means ± SE.

The present findings were in disagreement with that obtained by (Mahipal, et al. 2001) who showed that the variance due to GCA was more important for egg production and egg weight. In another study which reported by (Drumond, et al. 2014) on the layer performance of 4 strains of meat type quails using full diallel cross, resulted in a significant GCA for weekly egg number and egg yield.

Specific combining ability (SCA):

The SCA for egg number, egg weight and egg mass are presented in table (2). It can be observed that just EW trait resulted in a significant difference ($p \leq 0.05$) for SCA at the beginning of laying in the cross (W*Db), which means that Db line plays a role in laying large eggs because of its heavier BW, but Lb line as sire and dam reducing the egg weight in the next week.

Table (2). Specific combining ability for egg characteristics of crosses at first two weeks of laying period.

Period(wks.) Cross	Egg number		Egg weight		Egg mass	
	W1	W2	W1	W2	W1	W2
W*Lb	-0.12±0.01	-0.29±0.08	0.02±0.06ab	-0.01±0.02	-1.12±0.07	-3.11±0.68
W*Db	-0.04±0.07	0.08±0.01	0.23±0.11a	0.25±0.14	0.67±1.13	2.07±0.51
Lb*Db	0.47±0.47	0.31±0.37	-0.12±0.03b	-1.78±0.59	4.05±4.49	3.35±4.22
Sig.	Ns	Ns	*	Ns	Ns	Ns

Ns = Non-significant; *=significant at ($P \leq 0.05$); values = means \pm SE. Means having different letters within each column differ significantly.

However, the SCA values for egg number and egg mass of (Lb*Db) cross were higher and insignificantly positive at the first two weeks of laying period (0.47 & 0.31 eggs) and (4.05 & 3.35 gm), respectively. The present results are in disagreement with that found by (Mekky, et al. 2008) who mentioned that SCA was a significant source for egg production among crossbreed groups during all studied the ages.

Heterosis (H%):

Table (3) presented the heterosis percentage of egg number and egg weight for the crosses and its reciprocal crosses during the laying period. The results showed significant differences among all genotypes for both studied traits. In respect to EN characteristic, the reciprocal cross (Db*Lb) showed positive significant heterosis superiority by about (25% and 42%) for the interaction between first and second weeks of laying, respectively, which may reflect the Lb dam and Db sire.

Table (3). Heterosis percentage for egg number and egg weight of crosses and reciprocal crosses during the first two weeks of laying.

Period (wks.) Genotype		Egg number		Egg weight	
		W1	W2	W1	W2
Crosses	W*Lb	-27.02±0.68 ^b	-20.34±1.55 ^c	1.28±1.63 ^b	0.83±1.32 ^b
	W*Db	-0.77±7.81 ^{ab}	25.62±9.95 ^{ab}	8.09±2.07 ^a	6.8±1.16 ^a
	Lb*Db	-4.92±8.98 ^{ab}	11.24±6.87 ^{abc}	1.8±0.53 ^b	3.83±1.46 ^{ab}
Reciprocal Crosses	Lb*W	-8.18±18.37 ^{ab}	-13.91±15.25 ^c	1.1±0.69 ^b	2.73±1.19 ^{ab}
	Db*W	-25.97±7.24 ^b	0.61±13.39 ^{cb}	2.46±0.59 ^b	5.32±0.87 ^a
	Db*Lb	24.97±0.33 ^a	41.82±3.03 ^a	0.52±0.03 ^b	2.72±1.53 ^{ab}
Sig		*	**	**	*

*=significant at ($P \leq 0.05$); **=highly significant ($P \leq 0.01$); values = means \pm SE. Means having different letters within each column differ significantly.

Regarding to EW characteristic, the cross (W*Db) surpassed positively and significantly all other genotypes in the first week of laying (8.10) and didn't differ from its reciprocal (Db*W) in the second week of laying (6.8 and 5.3) respectively. The last result may mean that W line as sire interacted positively with the dam (Db) to produce heavier eggs. Also, its reciprocal cross affected by the sire (Db) in order to produce larger eggs, which reflect again the effect of Db line on the

egg size when crossed with W line. These results are in agreement with that reported by (Minvielle, et al. 2000) who indicated that the heterosis from crossing of two lines from different origins which were selected for early egg production affected significantly by the crossing. However the heterosis percentage for the egg number and egg weight depends on the age at the first egg and rate of the egg laying (Moritsu, et al. 1997; Sato, et al. 1989 and Piao, et al. 2002).

Reciprocal effect:

Table (4) presented the reciprocal effect on egg number, egg mass and egg weight during the first two weeks of laying periods. The results showed insignificant differences among the three genotypes in the first studied week for both egg

number and egg weight traits. While EP traits affected significantly ($p < 0.05$) by reciprocal cross in the second week. In respect to EN trait the reciprocal cross (Db*Lb) showed the highest positive significant value (0.65 egg) in the second week.

Table (4). Reciprocal effect on egg characteristics during first two weeks of laying.

Period (wks.)	Genotype	Reciprocal crosses			Sig.
		Lb*W	Db*W	Db*Lb	
Egg number	W1	0.48±0.48	-0.65±0.00	-0.35±0.40	Ns
	W2	0.18±0.45 ^{ab}	-0.6±0.12 ^b	0.65±0.20 ^a	*
Egg weight	W1	-0.07±0.07	-0.28±0.07	-0.08±0.01	Ns
	W2	0.05±0.03 ^a	-0.08±0.01 ^b	0.00 ^{ab}	*
Egg Mass	W1	4.55±4.51 ^a	-7.84±0.32 ^b	6.53±2.12 ^a	*
	W2	2.19±4.55 ^{ab}	-6.85±1.36 ^b	6.63±2.15 ^a	*

Ns = Non-significant; *=significant at ($p \leq 0.05$); values = means ± SE. Means having different letters within each row differ significantly.

Regarding to EW trait, the reciprocal cross (Lb*W) resulted in the highest value (0.05 gm) in the second week of laying. In respect to EM trait, it is significantly different among the genotypes in both studied weeks, in order to show the superiority for (Db*Lb) reciprocal cross, where it recorded the highest values (6.53 and 6.63 gm, respectively). While the reciprocal cross (Db*W) recorded the lowest values (-7.84 and -6.85 gm, respectively). Similar results were found by (Sharma, et al. 1992 and Mahipal, et al. 2001)

who showed significant reciprocal crosses effects on body weight, age at first egg and egg weight traits.

Maternal effect:

The M_E on egg number, egg mass and egg weight is presented in table (5) It can be observed that all studied characteristics differed significantly among the studied lines, and Db line resulted in the best values. In other words the Db dam plays a positive role in egg production traits.

Table (5). Maternal effect for egg characteristics during the first two weeks of laying.

Period(wks.) Line	Egg number		Egg weight		Egg mass	
	W1	W2	W1	W2	W1	W2
W	-0.12±0.32 ^b	-0.28±0.38 ^b	-0.23±0.002 ^c	-0.02±0.02 ^b	-2.19±3.22 ^b	-3.12±3.94 ^b
Lb	-0.55±0.05 ^b	-0.35±0.05 ^b	0.03±0.02 ^b	-0.07±0.02 ^b	-5.46±0.49 ^b	-3.94±0.48 ^b
Db	0.67±0.27 ^a	0.63±0.33 ^a	0.20±0.02 ^a	0.08±0.01 ^a	7.66±2.73 ^a	7.05±3.46 ^a
Sig.	*	*	**	**	*	*

*=significant at ($p \leq 0.05$); **=highly significant ($P \leq 0.01$); values = means ± SE. Means having different letters within each column differ significantly.

The present results are in disagreement with that found by (Barbato and Vasilatos Younken, 1991) who showed that maternal effects didn't change at earlier laying period. They added that the significant maternal effect at later ages may be due to the effect of endoplasmic inheritance which

plays a role in determination of specific maternal effect among lines. Another author mentioned that the differences between two strains for egg size or egg contents should not be the only source for maternal effect (Sabri, et al. 2000).

Genetic value:

Genetic values for egg characteristics during the first two week of laying are shown in table (6). The results showed that there were no significant effects for crosses on egg number and egg mass.

While egg weight appeared significant ($p < 0.05$) GV in the first week, where the (W*Db) cross resulted in the highest value (10.35 gm) followed by W*Lb cross (10.07 gm), and the lowest GV was recorded for the Lb*Db cross (9.82 gm).

Table (6). Genetic value for egg characteristics during the first two week of laying.

Period(wks.) Cross	Egg number/hen		Egg weight (gm)		Egg mass (gm)	
	W1	W2	W1	W2	W1	W2
W*Lb	4.24±0.62	4.38±0.33	10.07±0.15 ^{ab}	10.41±0.21	42.71±6.77	45.64±4.47
W*Db	4.52±0.65	5.36±0.45	10.35±0.15 ^a	10.81±0.06	46.82±5.81	57.81±4.38
Lb*Db	4.74±1.23	5.08±0.87	9.82±0.01 ^b	10.47±0.25	46.58±12.2	53.46±10.3
Sig.	Ns	Ns	*	Ns	Ns	Ns

Ns = Non-significant; *=significant at ($P \leq 0.05$); values = means ± SE. Means having different litters within each column differ significantly.

The last result indicates that Db dam play a positive role to increase the egg weight in its progeny that resulted from crossing it with W sire. Similar findings were obtained by (Razuki and AL-Shaheen, 2011) in chickens; they found that the genetic effects on egg number were insignificant, while it was significant on egg weight trait.

Heritability and genetic correlation:

The heritability, genetic correlation and phenotypic correlation coefficients within each studied line are represented in Table (7). In respect to heritability, which estimated directly from the relationship between parent and offspring, and indirectly from the random effects of both GCA and SCA. The results showed that the highest estimation values were recorded EN (0.53) and EW (0.51) in W line. All other estimations for other lines were moderate to small values. This result indicates that W line appears to have more additive genetic variance for the studied traits. Similar to the present results (Sato, et al. 1989)

reported heritability estimation for egg characteristic which had high values ranging from (0.62 to 0.84). Also (Stino, et al. 1982) mentioned that the heritability estimation obtained by regression method was high for some egg characteristic. Generally, (Berwary, et al. 2015) found that heritability estimation for EM was (0.57), in J. quail birds. While, (Hussen, et al. 2016) recorded low heritability estimate (0.21) for EW characteristic in brown J. quail birds.

Regarding to genetic correlation estimation (Table 7), the results show that the correlation coefficients between EW and EN were negative for both W and Db lines, but the coefficient (-0.59) in Db line was significant ($P < 0.05$), which reflect that Db line have a higher EW because it had higher body size, but W line had a smaller size.

Table (7). Heritability, genetic correlation and phenotypic correlation coefficients for some traits within studied quail lines.

Traits	Line	EN _{2weeks}	EW _{2weeks}
EN _{2weeks}	W	0.53	- 0.35
	Lb	0.28	-0.37
	Db	0.37	-0.4
EW _{2weeks}	W	-0.46	0.51
	Lb	NE	0.32
	Db	-0.59*	0.21

Heritability on the diagonal, phenotypic correlation above the diagonal and genetic correlation coefficients bellows the diagonal. *=significant at ($P < 0.05$); NE = Non-estimated.

The present results disagree with that found by (Mielenz, et al. 2006) in the Japanese quail, and also disagreed with the results that were reported by (Hidalgo, et al. 2011) who mentioned that the genetic correlations between EW and EN at different weeks of laying ranged from negative to positive values (-0.28 to 0.45).

In respect to phenotypic correlation coefficients (Table 7), all coefficients were negative and moderate to relatively small, which may due to the effect of both genetic and environmental factors. Moreover, (Choprakarn, et al. 1998) reported negative phenotypic correlation between the egg weight and few egg characteristics. On the contrary, (Kul and Seker, 2004) mentioned Positive correlation coefficients between some egg characteristics.

CONCLUSION

It is concluded from results of this research that, SCA and reciprocal cross have had significant effect ($p < 0.05$) among the studied genotypes for the studied traits, while the heterosis percentage for the same traits hadn't significant different ($p > 0.05$) among the studied genotypes, in particular, Db line might be used as dam in crossing for egg yield.

REFERENCES

- Amin EM, Kosba MA, El- Dlebs hany AE & El-Ngomy MA (2013). Heterosis, maternal and direct additive effects for growth traits in the Alexandria chickens. Egypt. Poult. Sci; 33: 1033-1051.
- Anang A, Mielenz N & Schuler L (2000). Genetic and phenotypic parameters for monthly egg production in White Leghorn Hens. J. Anim. Breed. Genet; 117: 407-415.
- Barbato GF & Vasilatos-Younken R (1991). Sex-linked and maternal effects on growth in chickens. Poult. Sci; 70: 709-718.
- Berwary MS, Oramari RA & Hussen SH (2015). Correlated response to selection for residual feed intake in Japanese quail (*Coturnix coturnix japonica*). Journal of University of Duhok; 18(1): Pp 135-144.
- Choprakarn K, Salangam I & Tanaka K (1998). Laying performance, egg characteristicistics and egg compositions in the indigenous hens. J. Natl. Res. Council. Thailand; 30:1-17.
- Dana N, Vander Waaij EH & Van Arendonk JM (2011). Genetic and phenotypic parameter estimates for body weights and egg production in Horro chicken of Ethiopia. Trop Animal Health Prod; 43: 21-28.
- Drumond ESC, Pires AV, Bonafe CM, Pinheiro SRF, Veloso RC, Amaral JM & Abreu LRA (2014). Egg laying and egg quality in meat type quails in diallel crosses. Genet. Mol. Res; 13 (4): 8118-8125.
- Drumond ESC, Pires AV, Veloso RC, Bonafe CM, Pereira IG, Costa LS, Abreu LRA (2015). Performance of meat type quails in diallel cross. Arq. Bras. Med. Vet. Zootec; 67 (1): 235-241.
- Duncan DB (1955). Multiple range and multiple F tests. Biometrics; 11: 1-42.
- Falconer DS (1988). Introduction to Quantitative Genetics. Third edition, John Wiley and Sons, New York, U.S.A.
- Hidalgo AM, Martins EN, Santos AL, Quadros TCO, Ton APS & Teixeira R (2011). Genetic characteristicization of egg weight, egg production and age at first egg in quails R. Bras. Zootec; 40: 95-99.
- Hussen SH, Hassan AM & AL-Khdri AMA (2016). Estimation of some genetic parameters for egg quality traits in Japanese quail (*Coturnix coturnix japonica*). Journal of University of Duhok; 19 (1): 32-37 (Special Issue).
- Isik F (2009). Analysis of Diallel Mating Designs. First edition, North Carolina State University, Raleigh, USA.
- Khaldari M, Pakdel A, Mehrbani Yeganeh H, Nejati Javaremi A & Berg P (2010). Response to selection and genetic parameters of body and carcass weights in Japanese quail selected for 4-week body weight. Poult. Scie; 89: 1834-1841.
- Koack C, Altan O & Akbas Y (1995). Japon bildircinlarının çeşitli verim özellikleri üzerine araştırmalar. Tr. J. of Vet. and Anim. Sci; 19: 65-71.
- Kul S & Seker I (2004). Phenotypic correlations between some external and internal egg quality traits in the Japanese quail (*Coturnix coturnix japonica*). International Journal of Poultry Science; 3:400-405.
- Lesson S & Summers JD (2005). Commercial Poultry Nutrition, 3rd edition, University books, P. O. Box. 1326, Guelph, Ontario, Canada ISBN 978-1-904761-78-5.
- Lotfi E, Zerehdaran S & Ahani Azari M (2012). Estimation of genetic parameters for egg production traits in Japanese quail (*Coturnix Coturnix japonica*). Arch. Geflügelk; 76 (2): 131- 135, 2012, ISSN 0003-9098.
- Luo PT, Yang RQ & Yang N (2007). Estimation of genetic parameters for cumulative egg numbers

- in a broiler dam line by using a random regression model. *Poult. Sci*; 86: 30-36.
- Mahipal RP, Venkateshwarlu YP, Prasad VL, Eshwaraiah K, Ravinder RV (2001). Combining ability analysis of a diallel cross involving three quail lines for production traits. *Indian Journal of Poult. Sci.* Volume: 36, Issue; 2: 147-152.
 - Mekky SS, Galal A, Zaky HI & Zein-EI-Dein A (2008). Diallel crossing analysis for body weight and egg production traits of two native Egyptian and two exotic chicken breeds. *Int. J. Poult. Sci*; 7: 64-71.
 - Mielenz N, Ronny R & Schuler L (2006). Estimation of additive and non-additive genetic variances of body weight, egg weight and egg production for quails (*Coturnix coturnix japonica*) with an animal model analysis. *Arch. Tierz*; 49: 300-307.
 - Minvielle F, Monvoisin JL, Costa J & Maeda Y (2000). Long-term egg production and heterosis in quail lines after within-line or reciprocal recurrent selection for high early egg production, *Brit. Poult. Sci*; 41:150-157.
 - Moritsu Y, Nestor KE, Noble DO, Anthony NB, Bacon WL (1997). Divergent selection for body weight and yolk precursor in *Coturnix coturnix japonica*: 12. Heterosis in reciprocal crosses between divergently selected lines, *Poult. Sci*; 76: 437-444.
 - Murakami AE & Arika J (1998). Produção de codornas japonesas. Jaboticabal: FUNEP, 79p (Report).
 - Piao J, Okamoto S, Kobayashi S, Wada Y & Maeda Y (2002). Study of heterosis effects on productive traits of Japanese quails: Heterosis effects on the crosses between large line and random bred population, *J. Poult. Sci*; 39: 139-146.
 - Piao J, Okamoto S, Kobayashi S, Wada Y & Maeda Y (2004). Purebred and crossbred performances from a Japanese quail line with very small body size. *Anim. Res*; 53: 145-153.
 - Razuki WM & AL-Shaheen SA (2011). Use of full diallel cross to estimate crossbreeding effects in laying chickens. *International Journal of Poult. Sci*; 10 (3): 197-204.
 - Saatci M, Omed H & Dewi IA (2006). Genetic parameters from univariate and bivariate analyses of egg and weight traits in Japanese quail. *Poult. Sci*; 85: 185-190.
 - Sabri HM, Khattab MS & Abdel-Ghany AM (2000). Genetic analysis for body weight traits of a diallel crossing involving Rhode Island Red, White Leghorn, Fayoumi and Dandarawi Chickens. *Annals of Agricultural Science Moshtohor*; 38: 1869-1883.
 - SAS Institute (2010). SAS User's Guide, Ver. 9.1: Statistics. SAS Institute Inc., Cary, NC.
 - Sato K, Fukuda H, Hediando YE & Ino T (1989). Heterosis for reproductive traits in reciprocal crosses of highly inbred lines of Japanese quail, *Jpn Poult. Sci*; 26: 70-73.
 - Sezer M (2007). Heritability of exterior egg quality traits in Japanese Quail. *Journal of Applied Biological Sciences*; 1 (2): 37-4.
 - Sharma AK, Johari D C, Kataria MC & Singh DP (1992). Combining ability analysis for egg production traits of light and heavy breed crosses of egg type chicken, *Indian Journal of Poult. Sci*; 27: 183-187.
 - Stino FKR, Kicka MA, Kamaer GA, Altakreti BTO (1982). Egg quality traits of the Japanese quail and their heritability in the subtropics. *Archiv für Geflügelkunde*; 3:104-108.
 - Williams, SM, Price SE & Siegel PB (2002). Heterosis of growth and reproductive traits in fowl. *Poult. Sci*; 81:1109-1112.

ههلسهنگاندنا بو ماوهی بو سبسکا ب ریکا لیکدانین دوورهی 2 - ساخاله تین هیکا

پوخته

قئ فکولینئ دا، 1307 جیجه لوک هاتنه ههلهتینان د دوو جارا ل دهوکئ، ژ سئ جوران: سپی، قههوی یئ قه بوی و قههوی یئ توخ، بژیئ نیک روژی ژ بو مه ره ما لیکدانئ. ل پروژی خودانکرنا په له وه را ل پشکا به ره مهئ گیانه وه ری ل کولیژا چاندنئ زانکویا دهوک، هه ریما کوردستانئ، عراق. نهف سالوخه تین هاتنه خواندن ل سهر هر دوو نقشا: ژبیئ گهه شتتا ره گه زی، ساخله تین به ره مهئینانا هیکا (کیشا هیکئ، ژمارا هیکا، باروستئ هیکئ)، ریژا ژ نافچونئ، ریژا پیتانن و ده رکه فتتا ژ هیکئ نهف پیقه رین بوماوهی هاتن ته خمین کرن. شیانا گشتیا لیکدانئ (GCA)، شیانا لیکدانا تایهت (SCA) کاریگریا لیکدانا باب و به ره بابا و هیزا دوو ره یئ (%H)، کاریگریا ماکئ (ME)، بهایئ خودانکرئ، و بهایئ لیکدانئ، بهایئ بوماوهی، شیانا قه گوهاستتا بوماوهی، پیقه گریدانا بوماوهی. گرنگرترین نهجامین هاتینه دیار کرن دقئ قه کولینئ دا دیار کر کو شیانین هاوکولکه بیین تابهت جیاوازین بهر چاف هه بون بو سالوخه تین به ره مهئ هیکا. چ جیاوازین بهر چاف نه بون د ناف بهرا لاینن قه کولینئ د هیزا دووره گ بو سالوخه تین سهنگا هیکئ و ژمارا هیکا. بهلئ کارتیکرنا دوورهی لسره وه کهه فی بهروفاژی یا بهرچاف بو. دیسان چ جیاوازین بهرچاف نه بون د ناف بهرا لاینن قه کولینئ دا و کارتیکرنا بهه یئ پهروه دهی دا د هه می ماوی قه کولینئ دا. نهجامئ قه کولینئ دیار دکن کو لاینن قهوائی قه بوی وهک باب بو به ره مه ئینانا به ره مهئ هیکا.

التقييم الوراثية للسمان باستخدام التهجين ثنائي الأليل الكامل 2- إنتاج البيض

الخلاصة

تم فقس عدد 1307 صوصاً على دفعتين من ثلاثة خطوط للسمان الياباني (الأبيض والبني الفاتح والبني الغانق). أجريت التجربة في مزرعة الدواجن بقسم الإنتاج الحيواني في كلية الزراعة- جامعة دهوك، إقليم كردستان، العراق. هدفت التجربة لتقييم الأساس الوراثي لثلاثة خطوط من السمان الياباني لصفات إنتاج البيض (وزن البيض و عددها وكتلتها) باستخدام التهجين ثنائي الأليل الكامل. وقد درست المعالم الوراثية التالية: قدرتي التوافق العامة والخاصة والتأثير الأمي والتهجين التبادلي وقوة الهجين، بالإضافة لتقدير القيمة الوراثية ومعامل التوريث لصفتي إنتاج البيض (وزن البيض و عددها وكتلتها) لم تختلف معنوياً. سجلت قدرة التوافق الخاصة فروقا معنوية لصفات إنتاج البيض. لم تختلف التراكيب الوراثية المدروسة فيما بينها معنوياً لظاهرة قوة الهجين بالنسبة وزن البيض و عدده. لكن تأثير الهجين العكسي كان معنوياً على خصائص إنتاج البيض. لم يكن التأثير الأمي معنوياً في التأثير على خصائص إنتاج البيض. وقد استنتج الى انه يمكن استخدام الخط البني الغامض كامهات في حال إنتاج البيض.