# IMPROVING THE PRODUCTIVITY AND REPRODUCTIVITY OF BAHEIJ CHICKENS THROUGH CROSSING

By

Effect of upgrading on: Growth traits, chicks viability, body conformation and carcass traits

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Abstract: Crossing trials were carried out to evaluate the effect of upgrading (increasing blood of Silver Montazah, SM chickens) on certain traits (body weight, growth rate and growth efficiency percentages, birds viability percentages at different ages) throughout crossing selected Montazah males for mature body weight as a parent line with Baheij (Bj) dams producing three generations progenies [(the 1<sup>st</sup> generation (½ SM + ½ Bj), the 2<sup>nd</sup> generation (¾ SM + ¼ Bj) and the 3<sup>rd</sup> generation (7/8 SM + 1/8 Bj)]. Body measurements (BM) and body conformation (BC) at 8 and 12 wks of age and carcass traits at 16 and 20 wks of age were studied. Results were as follows:

- 1- Averages body weight (BW) of upgraded group chicks surpassed (P<0.001) those of Bj at all studied ages. Moreover, chicks of the  $3^{rd}$  generation were the heaviest (P<0.001) compared to those of the other generations.
- 2- Upgrading improved (P<0.001) growth rate percentage (GR%) at 0-16 wks of age while the contrary was found at the period 8-12 wks of age. The 3<sup>rd</sup> generation had the highest values of GR% from hatch to 16 wks and 12-16 wks of age.
- 3- Averages growth efficiency (GE) of upgraded groups surpassed (P<0.001) the purebred at 0-4, 0-12, 0-16 and 0-20 wks of age. the 3<sup>rd</sup> generation) had the highest (P<001) averages GE at 4-8, 0-12, 12-16, 0-16 and 16-20 wks of age. Moreover, Males had the higher averages concerning studied growth traits compared to the females.
- 4- Only generation affected viability (P<0.01) of chicks through the period 4-8 wks of age.

- 5- Averages of BW, shank length (ShL) and keel length (KL) of birds increased (P<0.01) by age. Also, the birds of the 3<sup>rd</sup> generation were heavier (P<0.05) than those of the purebred. On the other hand, upgrading did not improve any of the body conformation estimated values.
- 6- Both of generation and genotype significantly affected carcass and giblets weights where the averages of the 3<sup>rd</sup> generation were higher than those of the 2<sup>rd</sup> generation and the birds of upgraded group had higher averages for all traits studied compared to the purebred group. Also, the different between averages of most studied traits at 16 and 20 wks of age was highly significant.

It could be concluded that upgrading through crossing SM sires to Bj dams improved growth traits, chick viability, certain body conformation and carcass traits of progeny of both F2 and F3.

#### INTRODUCTION

Study of Abdalla (2007) showed that Baheij breed reached genetic equilibrium. Therefore, upgrading was suggested to overcome this problem to improve the productivity of this strain. Baheij breed was developed (Mahmoud et al., 1979) through crossing with Silver Montazah strain (Mahmoud et al., 1974). Therefor, crossing Silver Montazah males to Baheij females had to be applied in order to overcome the problem of reaching genetic equilibrium in Baheij breed, thus allowed effective selection for certain productive traits.

Crossing was found to be effective for improving body weight as reported by Shebl et al. (1995), Nawar et al. (2004), and Amin (2007) who found positive heterosis in body weight at 6 and 12 wks of age in both sexes, also, Nestor et al. (2006) and Amin (2009) reported positive effect of backcrossing on growth traits in turkey. Moreover, Mostafa and Nofal (2000), Amin (2008 and 2009) found significant difference between the two sexes in body weight at different ages.

Crossbreeding was found to improve chick viability (El-Turkey, 1981; Nawar and Abdou, 1999; Nawar et al., 2004).

Several reports have demonstrated that crossbreeding seems to improve body measurements (Abou-El-Ella, 1982 and Sharaf et al., 2006) and body conformation (El-Turkey, 1981), while no significant differences were found between purebred and crossbred as reported by the same auther.

Several investigators found positive effect of crossing on edible meat or giblets percentages (Ali, 1979; Khar, 1981; and Mandour et al., 1996),

while others found no heterotic effect of crossing on the same traits (El-Turkey, 1981; Sharaf et al., 2006). Moreover, pure parents were a significantly source of variation for body weight and all the carcass traits (Mandour et al., 1996).

The aim of the present study is to find out the effect of upgrading on body weight at different ages, growth rat, growth efficiency and birds viability percentages at different periods. Also, body conformation at 8 and 12 wks of age and carcass traits at 16 and 20 wks of age were studied.

## MATERIALS AND METHODS

This study was conducted at El-Sabahia Poultry Research Station, Alexandria, Animal Production Research Institute, Agriculture Research Center, Egypt.

# Breeding and management:

# **Upgrading Line:**

In the first generation, females of the Baheij (Bj) breed were crossed to selected Silver Montazah (SM) males as a parent line according to the individual mature BW (average), thus, the females produced (1/2SM + 1/2Bj) were backcrossed to the same parent line throughout two additional generations [produced (3/4SM + 1/4Bj) and (7/8SM + 1/8Bj), in the two generations, respectively].

#### Control line:

Chicks of Baheij breed were randomly chosen to establish a pedigreed control population.

Continuous lighting was provided from hatching to 8 wk of age, at that time, the photoperiod was reduced to 12 h/d. and remained at this level during the rearing period. After the rearing period, at 20 weeks of age, females were housed in breeding pens (10 pullets + 1 male, each). The birds were fed a starter diet (19% crude protein and 2800 Kcal) up to 8 weeks of age, grower diet (15% crud protein and 2700 K cal) up to 20 weeks (17% crude protein and 2850 Kcal). Thereafter, feed and water were supplied ad libitum. The average number of progeny reared in the three generations was 68, 135, 133 chick for the BjBj line, and 210, 315, 329 chick for the SM x Bj genotype, respectively.

#### The studied traits:

- Body weights (BW) at hatch, 4, 8 (not tabulated and presented in Figures 1 and 2). Also, BW 12, 16, and 20 wks of age presented in Tables.

- Growth rate using the following equation (Lerner and Asmundson, 1932).

$$GR = [W_2 - W_1/1/2(W_2 + W_1)] \times 100$$

Where: GR: rate of growth, W1: the initial weight,

W<sub>2</sub>: the second weight.

- Growth efficiency through different periods of growth using the following equation (Gondwe and Wollny, 2005):

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Where, GE is growth efficiency per time period =  $t_i$ ; WG<sub>ti</sub> is weight gain at time =  $t_i$ ; LW<sub>to</sub> is live weight at time =  $t_0$ 

- Chick viability through the growth period (0-8 weeks) of age.
- Body measurements:

Shank length, keel length and breast width which was measured with a modified vurger angle meter in millimeters. It was measured at a point very near to the fronh end of the breast bone and ¾ inch down from the keel. at 8 and 12 weeks of age at the 3<sup>rd</sup> generation, and body conformation was estimated using the following equations (Nordskog, 1976, Mahmoud *et al.*, 1980 and El-Turkey, 1981):

 $C_1 = \sqrt[3]{BW}$  / shank length, gm/cm

C<sub>2</sub> = BW/shank length, gm/cm

 $C_3 = \sqrt[3]{BW}$  / keel length, gm/cm

 $C_4 = BW / keel length, gm/cm$ .

- Random sample of 6 cocks at 16 and 20 weeks of age from each genotype for the two generations were used to study the carcass traits (Absolute values and percentages of carcass, legs, gizzard liver and heart. Statistical analysis:

Data of growth traits were analyzed using fixed models SAS institute (1988):

$$Y_{ijkl} = U + Gn_i + Gt_j + S_k + (GnGt)_{ij} + (GnS)_{ik} + (GtS)_{jk} + GnGtS)_{ijk} + e_{ijkl}$$

Where:  $Y_{ijkl}$ = an observations, U = overall mean,  $Gn_i$  = the fixed effect of  $i^{th}$  generation,  $Gt_j$  = the fixed effect of  $j^{th}$  genotype,  $S_k$  = the fixed effect of  $k^{th}$  sex, and  $(GnGt)_{ij}$ ,  $(GnS)_{ik}$ ,  $(GtS)_{jk}$  and  $(GnGtS)_{ijk}$  = effects of the interactions between the three factors studied, and e iikl = random error.

The other traits which studied were analyzed using fixed models SAS institute (1988):

$$Y_{ijk} = U + Gn_i + Gt_j + (GnGt)_{ij} + e_{ijk}$$

Where:  $Y_{ijk}$ = an observations, U = overall mean,  $Gn_i$ = the fixed effect of  $i^{th}$  generation,  $Gt_j$ = the fixed effect of  $j^{th}$  genotype,  $(GnGt)_{ij}$ = effect of the interaction between the two main factors, and  $e_{ijk}$ = random error. Significant differences among means were tested by Duncan Test (1955).

#### RESULTS AND DISCUSSION

### 1 -Body weight (BW):

Least square means for BW as affected by generation, genetic group, for males and females are presented in Table 1 and Figures 1 and 2. Concerning the effect of generation, the results showed that, chicks of the 3<sup>rd</sup> generation were the heaviest (P<0.001) compared to those of the 1<sup>st</sup> and 2<sup>nd</sup> generations (769.1, 1031.2 and 1299.5 gm) at 12, 16 and 20 wks of age, respectively. The same trend was found at the early periods of growth (0, 4, and 8 wks of age as shown in Figure 1). On the other hand, while the averages of body weight of the 1<sup>st</sup> and 2<sup>nd</sup> generations were nearly similar at 12 wks of age, the superiority of the 1<sup>st</sup> one over the 2<sup>nd</sup> generation (P<0.001) was found at 16 wks of age while the opposite was found at 20 wks of age.

Results showed that the least squares means of body weight of upgraded group surpassed (P<0.001) those of the contro at all ages studied. The positive effect of crossing agrees with the finding of Yalcin *et al.* (2000), Nawar *et al.* (2004), Mohamed (2003), Aly *et al.* (2005) and Amin (2007) on chicken and Amin (2009) on turkey. Nestor *et al.* (2006) reported that for maximum gains per generation, backcrossing probably should be used for maximum of two or three generations.

Males were heavier (P<0.001) than females at all ages studied except those of 1 day of hatch which had nearly similar weight averages. Same results were reported by Mostafa and Nofal (2000) and Amin (2008). In contrary, Nestor *et al.* (2006) found that males in the F1 generation did not differ from expected for body weight at any age but the females of that cross had higher body weight than expected at 16 and 20 weeks of age.

Statistical analysis revealed no significant interaction between the three main factors except in BW at 8, 16 and 20 wk of age. Significant (P<0.001) effect of interaction between generation (Gn) x genotype (Gt)

was found. The birds in the  $3^{rd}$  generation (7/8 SM + 1/8 Bj) were the heaviest (1171.3 and 1418.5g) at 16 and 20 weeks of age, respectively.

# 2-Growth rate percentage (GR%):

Generation affected (P<0.001) GR% at 4-8 and 8-12 wk of age (Figure 3). The results in Table 2 showed that least squares means of GR% of birds differed (P<0.001) through the three consecutive generations, where birds of the 3<sup>rd</sup> generation had the highest values of GR% through the intervals 12-16, and 0-16 wks of age (28.6 and 186.2%, respectively), while the 2<sup>nd</sup> generation had the highest value at 16-20 wks of age (24.4%). On the other hand, upgrading improved (P<0.001) GR% at 0-16 wk of age (186% vs. 184.6%) for both upgraded and purebred, respectively. Also, upgrading slightly increased GR% values at the 1<sup>st</sup> periods except at 8-12 wks of age. The contrary was found where the GR% of the purebred was higher (P<0.001) than that of the upgraded group (Figure 4). Results of some investigators were in harmony with the results in this study. Shebl et al. (1995) reported that the crossbreds grew significantly faster than the purebred from the hatch day to 8 wks of age, but vice versa at 8 to 12 wks of age also, Nawar et al., (2004) reported that crossbreeding improved significantly growth rate during early intervals of age (4-6 wks).

Males had significantly (P<0.001) higher values of GR% at the periods 0-4, 0-16, and 0-20 wks of age than females as shown in Figures 3, 4 and Table 2. These results were in agreement with thaose reported by Mostafa and Nofal (2000) and Amin (2008 and 2009) who found significant differences between both sexes in body weight. All interactions between the main factors were not significant except that for GR% at 0-16 wks of age. The birds of upgraded groups at both the 1<sup>st</sup> and 3<sup>rd</sup> generations were equal and had significant GR% (185.5%).

#### 3- Growth efficiency (GE):

Figure 5 and Table 3 shows that generation affected GE significantly (P<0.001) at 4-8 wks of age. Also, the same effect was found at 0-12 (P<0.05) and at 12-16, 16-20, 0-16 and during the whole period (0-20) (P<0.001) where the birds of the 3<sup>rd</sup> generation had the highest means of GE at all the mentioned periods except at 16-20 where the birds of the 2<sup>nd</sup> generation had the highest values (21.76, 0.34, 0.21, 28.14 and 38.77, respectively, Table 3). In addition, results show that upgrading improved significantly GE at 0-4, 0-12, 0-16 and 0-20 wks of age (P<0.001) (Figure 6

and Table 3). Moreover, Baheij males had significantly higher values of GE at the most periods studied (P<0.01 to P<0.001).

On the other hand, results revealed highly significant effects of the interaction between Gn x Gt for GE at 0-12, 12-16, 0-16 and 0-20 wks of age. The genotype 7/8SM + 1/8Bj (at the  $3^{rd}$  generation) had the best growth efficiency. Amin (2009) found that all crosses and backcrosses in turkey had significantly the lowest means of growth efficiency through the period (4-20 wks of age) concerning the three generations studied on turkey. On the other hand, he found that the difference between overall means of growth efficiency for both sexes was significant at all periods studied except for 12-16 and 12-20 wks of age.

# 4- Body measurements (BM) and body conformation (BC):

Results presented in Table 4 suggested that age affected body weight, shank length and keel length significantly (p<0.01) where the averages values of these traits at 12 wks of age were higher than those at 8 wks of age. Also, BW averages differed significantly in both genotypes (upgraded and control) where birds of 7/8 SM + 1/8 Bi were heavier (P<0.05) than those of pure Bj (663.3 vs 494.4 g). Moreover, males were heavier (P<0.01) than females (733.1 vs 558.5). On the other hand, it seemed that upgrading did not improve BC, estimated by any of  $C_1$ ,  $C_2$ ,  $C_3$ or C<sub>4</sub>. Comparing body conformation for males and females, it was found that C2 and C4 were more applicable (P<0.01) than C1 and C3. On the other hand, it was found that C3 is not accurate measurement for body conformation at 8 and 12 wks of age compared to C1, C2 and C4 where they were considered more refined (P<0.01). There was significant interaction between age x generation for shank length (P<0.01), keel length and BC values (P<0.05), while interaction effects were not significant for BW or breast width.

These results were in agreement with those reported by El-Turkey (1981) who reported that differences were not significant among body measurements of purebreds and crossbreds at certain ages (8 and 12 wks), while Abou-El-Ella (1982) reported that crossbreeding was found to be effective on body measurements, particularly on keel length and breast width. In addition, El-Turkey (1981) reported that crossbreeding seems to improve body conformation (C<sub>1</sub>, C<sub>2</sub>, C<sub>3</sub> and C<sub>4</sub>). In contrary, Nestor (1971) found that crosses of large strains did not show any heterotic effects in body conformation in turkey.

On the other hand, our results disagreed with those reported by Sharaf et al. (2006) concerning both shank and keel length at 4 weeks in quail while the results of the same authers concerning both traits showed that average of reciprocal crossbreds surpassed both of pure and crossbred averages at 5 wks of age. General increase in some body measurements in each genotype as age increased and this finding in agreement with Adedeji et al. (2006).

# 5-Viability percentage (V%):

Results in Table 5 showed that viability of chicks differed significantly (P<0.01) through the period 4-8 wks of age for the three generations studied, while there were no significant differences between the two genotypes or between the two sexes at all the periods studied. Also, all the interactions between the three main factors had no significant effects on this trait in the different periods studied. The results of this disagreed with those reported for some investigators who confirmed the superiority of crossbred over purebreds in viability (Fairful, 1990; Mandour et al., 1992; Aly et al., 2005; Amin, 2007).

# 6-Carcass traits:

Results in Table 6 showed that live body weight, and carcass values differed significantly in birds of the 3<sup>rd</sup> generation compared to that of the 2<sup>nd</sup> generation at 16 and 20 wks of age (1278g, 907.1g and 69.4% vs. 1119g, 755.7g and 67.5%, respectively). Also, gizzard, and heart weights and legs percentage differed significantly in the two generations where the values of the 3<sup>rd</sup> generation were higher than that of the 2<sup>nd</sup> generation.

On the other hand, it was found that genotype affected (P<0.01) live body weight, carcass weight, where the averages of these traits were higher in upgraded group than those of pure Baheij birds. Also, gizzard and heart weights, and both the legs weight and percentage were affected (P<0.01) by the same trait. Age affected (P<0.01) live body weight, legs weight, carcass, liver and heart weights and percentages, where the values of the birds of 20 wks of age were higher than those of 16 wks except gizzard percentage where the contrary was found (P<0.01).

There were significant effects of all types of interaction on some carcass traits (Table 6). Comparisons of upgraded group relative to pure Baheij chickens were associated with superior effects of upgrading on most live body weight and carcass traits compared to Baheij group.

These results were in agreement with those reported by several investigators, that crossbreds were superior to purebreds concerning edible meat (Ali, 1979; Khar, 1981). In addition, Mandour *et al.* (1996) reported that the overall mean of edible giblets percentages were greater (P<0.05) for Silver Montazah line crosses than corresponding pure strain.

It could be concluded that crossing Baheij chickens (as dams) with Silver Montazah (as a sire parent) throughout three generations (upgrading) improved growth traits (body weight, growth rate percentage and growth efficiency) of progeny of both F2 and F3. These results are in agreement with several authors (Emmerson et al., 1991; Ye et al., 1997; Mostafa and Nofal, 2000) in chicken and Amin (2009) in turkey.

# **CONCLUSION**

Comparing the effect of upgrading with that of the purebred (Baheij), it was found that upgrading had more influence on average body weight at 20 wks. These held true for body weight of purebred and crossbred at different ages (4, 8, 12 and 16 wks of age).

Further studies should be made to cross the purebred Bj sires to 7/8SM, 3/4 SM and 1/2 SM at different generations to reach 7/8 Bj and 15/16 Bj. The last mentioned cross will be compared to selected Bj to investigate the improvement concerning upgrading, either-way (increasing the blood of Silver Montazah. or increasing the blood of Bj strain) in order to overcome the problem of genetic equilibrium in Bj chickens.

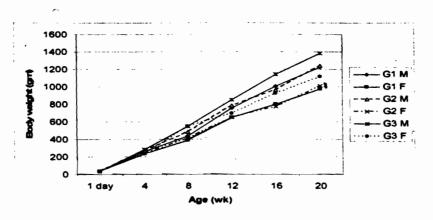


Figure 1. Effect of generation on body weight for males and females

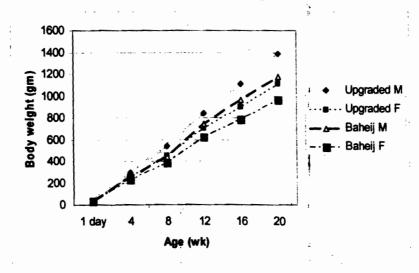


Figure 2. Effect of genotype on body weight for males and females

**Table (1):** Effect of genotype and generation on least squares means and  $\pm$  standard error of body weight of male and female birds at different ages studied

		Body weight, g										
Generation	Genotype	12 wks				16 wks		20wks				
		Male	Female	Average	Male	Female	Average	Male	Female	Average		
	1/2SM+1/2Bj	769.1 <u>+</u> 12.0	687.9±14.3	728.5±9.3	1041.9±16.5	840.1±19.8	941.0 <u>+</u> 12.9	1264.6+22.2	1028.8+26.4	1146.7+17.2		
	BjBj	733.0 <u>+</u> 20.8	611.4±20.3	672.2±14.5	974.9 <u>+</u> 28.4	764.4 <u>+</u> 28.8	869.7±.20.2	1181.9 <u>+</u> 37.9	921.9 <u>+</u> 38.9	1051.9 <u>+</u> 27.2		
Overall mean		751.1±12.0	649.6+12.4	700.3±8.6 <sup>8</sup>	1008.4+16.8	802.3+17.5	905.4+12.0 <sup>8</sup>	1223.2+21.9	975.4+23.5	1099.3+16.1 <sup>C</sup>		
,	3/4SM+1/4Bj	818.7±10.4	665.5±10.7	742.1 <u>+</u> 7.4	986.3±14.2	814.1±14.7	900.2±10.2	1306.9+18.4	1058.4+19.3	1182.6+13.4		
	BjBj	750.6±17.2	627.2±15.2	688.9±11.5	952.9+23.5	779.0+21.1	866.0+15.8	1177.8+30.5	970.25+27.4	1074.0+20.7		
Ove	rall mean	784.7 <u>+</u> 10.1	646.4+9.3	715.5+6.8	969.6+13.7	769.5+12.9	883.1 <u>+</u> 5.4 <sup>C</sup>	·1242.3+17.8	1014.3 <u>+</u> 16.8	1128.32+12.2 <sup>8</sup>		
1 2 1	7/8SM+1/8Bj	943.8+8.2	757.0±10.1	850.4±6.5	986.3±14.2	1032.2+14.0	1171.3+8.0	1585.7±14.6	1251.3±18.3	1418.5±11.7		
	BjBj	748.9 <u>+</u> 35.1	626.9±30.7	687.9+23.3	952.9±23.5	811.2+41.9	891.0±31.8	1175.4±26.3	985.6+56.5	1080.5±41.8		
Ove	rall mean	846.3±18.0	691.9+16.1	769.1±12.1 <sup>A</sup>	1140.6+24.6	922.0+22.1	1031.2+16.5 <sup>A</sup>	1380.6+31.9	1118.5+29.34	1299.5+21.7 <sup>A</sup>		
	of upgraded groups	843.9±6.0	703.5±6.8	773.7±4.5°	1112.9+8.2	895.5±9.5	1004.2±6.3ª	1385.7±10.8	1112.84+12.5	1249.3±8.2ª		
	m of Baheij breed	744.2 <u>+</u> 14.8	621.8±13.3	682.5±9.9 <sup>b</sup>	966.2±20.2	784.9±18.4	875.5±13.6 <sup>b</sup>	1178.4+26.3	959.3 <u>+</u> 24.5	1068.8±18.0 <sup>b</sup>		
Ove	rall mean	744.0±8.0°	662.6+7.5	773.9	1039.56+10.9°	840.2+10.3°	1011.6	1282.0+14.2°	1036.1±13.87	1256.3		

SM: Silver Montazah strain and Bj: Baheij breed,

- Means having different letters in every column and within every factor of treatments are significantly different (P<0.05),

- All main factors studied had highly significant (0.001) effect on the body weight while all interactions between them were not significant except that for body weight at 16 and 20 wks of age, which were influenced (p<0.001) by the interaction of generation x genotype.

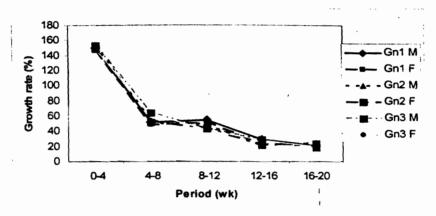


Figure 3. Effect of generation on growth rate percentage for males and females

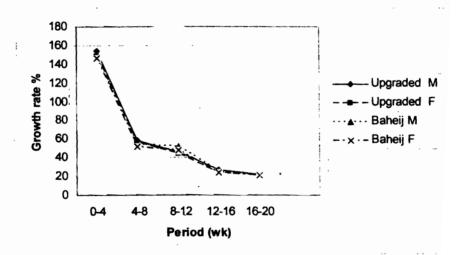


Figure 4. Effect of genotype on growth rate percentage for males and females

Table (2): Effect of genotype and generation on least squares means and ± standard error of growth rate percentage of male and female birds at different ages studied

G	Genotype	0-12 v	vks (Growth	period)		12-16 wks			16-20 wks			
Generation		Male	Female	Average	Male	Female	Average	Male	Female	Average		
•	1/2SM+1/2Bj	180.9±0.6	180.4±0.7	180.7±0.4	30.1±1.0	20.6 <u>+</u> 1.2	25.33±0.8	20.3+0.9	21.3 <u>+</u> 1.1	20.8±0.7		
' '	BjBj	181.7 <u>+</u> 0.9	178.3±0.5	180.0±0.7	28.5±1.8	24.1±1.8	26.3 <u>+</u> 1.3	20.1±1.6	22.6 <u>+</u> 1.6	21.4 <u>+</u> 1.1		
Overs	il mean	181.3 <u>+</u> 0.6	179.4±0.6	180.3 <u>+</u> 0.4	29.3 <u>+</u> 1.0	22.3±1.1	25.8±0.8 <sup>8</sup>	20.2 <u>+</u> 0.9	22.0 <u>+</u> 1.0	21.1 <u>+</u> 0.7 <sup>8</sup>		
2	3/4SM+1/4Bj	183.4±0.5	179.9±0.5	181.6 <u>+</u> 0.4	19.1 <u>+</u> 0.9	21.2+0.5	20.12 <u>+</u> 0.7	28.1 <u>+</u> 0.8	25.5±0.8	26.8 <u>+</u> 0.5		
2	BjBj	182.4+0.8	179.2±0.8	180.8 <u>+</u> 0.5	23.4 <u>+</u> 1.5	22.3±1.3	22.9±0.9	21.7 <u>+</u> 1.3	22.2 <u>+</u> 1.1	22.0 <u>+</u> 0.8		
Overs	il mean	182.9 <u>+</u> 0.5	179.5+0.4	181.2 <u>+</u> 0.3	21.3 <u>+</u> 0.9	21.7 <u>+</u> 0.8	21.5 <u>+</u> 0.6 <sup>C</sup>	24.9 <u>+</u> 0.7	23.9 <u>+</u> 0.7	24.4 <u>+</u> 0.5 <sup>A</sup>		
,	7/8SM+1/8Bj	185.1±0.4	181.2±0.5	183.1 <u>+</u> 0.3	32.0±0.7	30.1 <u>+</u> 0.9	31.0±0.6	18.5±0.6	19.0 <u>+</u> 0.8	18.8 <u>+</u> 0.5		
3	BjBj	181.7±1.6	179.1±1.4	180.4 <u>+</u> 1.1	26.6 <u>+</u> 2.9	25.7 <u>+</u> 2.6	26.2 <u>+</u> 2.0	19.2 <u>+</u> 2.5	19.7 <u>+</u> 2.3	·19.4 <u>+</u> 1.7		
Overs	ll mean	183.4 <u>+</u> 0.8	180.1 <u>+</u> 0.8	181.7 <u>+</u> 0.6	29.3+1.5	27.9±1.4	28.6 <u>+</u> 1.0 <sup>A</sup>	18.9+1.3	19.3 <u>+</u> 1.2	19.1 <u>+</u> 0.9 <sup>C</sup>		
	n of upgraded oups	183.1.3	180.5±0.3	181.8±0.2	27.1 <u>+</u> 0.5	23.9 <u>+</u> 0.6	25.5±0.4	22.3 <u>+</u> 0.4	21.9 <u>+</u> 0.5	22.1 <u>+</u> 0.3		
	ean of Baheij reed	181.9±0.7	178.8 <u>+</u> 0.6	180.4 <u>+</u> 0.5	26.2±1.3	24.0 <u>+</u> 1.1	25.1 <u>+</u> 0.8	20.4 <u>+</u> 1.1	21.5 <u>+</u> 1.0	20.9 <u>+</u> 0.7		
Overs	il mean	182.5±0.4	179.7+0.4		26.6±0.7	24.0±0.6		21.3+0.6	21.7+0.6			
	•			Signi	ficance of:							
Gene	eration		NS		•	**		***				
Gen	otype		NS			N		NS				
	Sex		NS			N	S		NS			

# Cont. Table (2)

			Growth rate percentage										
Generation	Genotype	12-20	wks (Rearing	period)		0-16 wks		0-20 wks					
		Male	Female	Average	Male	Female	Average	Male	Female	Average			
. 1	1/2SM+1/2Bj	50.8 <u>+</u> 1.4	41.3 <u>+</u> 1.6	46.0 <u>+</u> 1.1	186.9±0.2	184.0 ±1.3	185.5±0.2	187.5 ±0.6	186.9±0.7	187.2 <u>+</u> 0.4			
` <b>'</b>	BjBj	46.5 <u>+</u> 2.3	45.8 <u>+</u> 2.4~	~46.1 <u>+</u> 1.7	186.2 <u>+</u> 0.4	182.7 <u>+</u> 0.4	*184.4±0.3	188.6 <u>+</u> 1.9	185.8±0.9	187.2 <u>+</u> 0.7			
Over	rail mean	48.6 <u>+</u> 1.4	43.5 <u>+</u> 1.9	46.1±1.0	186.5±0.2	183.3 <u>+</u> 0.2	184.9±0.2 <sup>B</sup>	188.1 <u>+</u> 0.6	186.3 <u>+</u> 0.6	187.2±0.4			
2	3/4SM+1/4Bj	46.4±1.1	45.6±1.2	46.0 <u>+</u> 0.8	186.2 <u>+</u> 0.1	183.6 <u>+</u> 0.2	184.9±0.1	189.6±0.5	187.2±0.5	188.4 <u>+</u> 0.3			
4	BjBj	44.6 <u>+</u> 1.9	43.8 <u>+</u> 1.7	44.2 <u>+</u> 1.3	185.9 <u>+</u> 0.3	183.2 <u>+</u> 0.3	184.6±0.2	188.7 <u>+</u> 0.8	186.5±0.7	187.6±0.5			
Overall mean		45.5±1.1	44.7±1.0	45.1±0.8	186.1±0.2	183.4 <u>+</u> 0.2	184.7 <u>+</u> 0.7 <sup>8</sup>	189.1±0.5	186.9±0.4	186.9 <u>+</u> 0.4			
3	7/8SM+1/8Bj	49.7 <u>+</u> 0.9	48.3±1.1	49.0 <u>+</u> 0.7	186.9 <u>+</u> 0.2	184.0 <u>+</u> 0.3	185.5±0.1	187.5±0.6	186.9 <u>+</u> 0.7	187.2 <u>+</u> 0.4			
	BjBj	45.2 <u>+</u> 3.9	45.7±3.5	45.4 <u>+</u> 2.6	186.2 <u>+</u> 0.4	182.7 <u>+</u> 0.4	184.4 <u>+</u> 0.3	188.6 <u>+</u> 0.9	185.8 <u>+</u> 0.9	187.2 <u>±</u> 0.7			
Over	rall mean	47.4 <u>+</u> 2.0	47.0±1.8	47.2 <u>+</u> 1.3	187.5±0.3	184.9±0.3	186.2±0.2 <sup>A</sup>	189.6 <u>+</u> 0.8	187.4 <u>+</u> 0.8	188.5±0.6			
	all mean of ded groups	49.0 <u>+</u> 0.7	45.0 <u>+</u> 0.8	47.0 <u>+</u> 0.5	187.4 <u>+</u> 0.1	184.5 <u>+</u> 0.1	186.0 <u>+</u> 0.1ª	189.3 <u>+</u> 0.3	187.5 <u>+</u> 0.3	188.4 <u>+</u> 0.2			
	nean of Baheij breed	45.4 <u>+</u> 1.6	45.1 <u>+</u> 1.5	45.3 <u>+</u> 1.1	186.0 <u>+</u> 0.3	183.2 <u>+</u> 0.2	184.6±0.2b	188.6 <u>+</u> 0.7	186.2 <u>+</u> 0.6	187.4 <u>+</u> 0.4			
Over	all mean	47.2 <u>+</u> 0.9	45.1±0.9		186.7±0.1 <sup>X</sup>	183.9±0.1		188.9±0.4 <sup>X</sup>	186.8±0.3 <sup>Y</sup>				
					Significance	e of:							
Ge	neration		NS						NS				
Ge	enotype		NS			***			NS				
Sex			NS			***			***				

SM: Silver Montazah strain and Bj: Baheij breed,
- Means having different letters in every column and within every factor of treatments are significantly different (P<0.05),
- All interactions between the main factors were not significant except that for growth rate % at 0-16 period, which was influenced (p<0.001) by the interaction of generation x genotype.

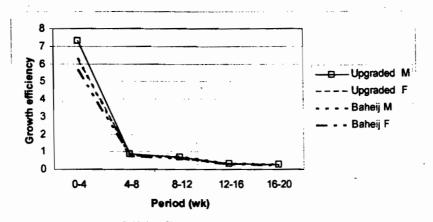


Figure 5. Effect of generation on growth efficiency for males and females

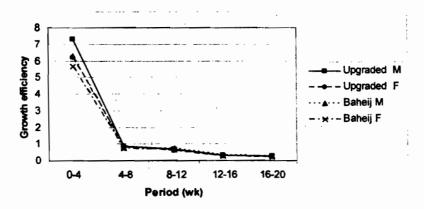


Figure 6. Effect of genotype on growth efficiency for males and females

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**Table (3):** Effect of genotype and generation on least squares means and  $\pm$  standard error of growth efficiency of male and female birds at different periods studied

Comono						Growth effic	eiency					
Genera tion	Genotype	12-20	wks (Rearing	period)		0-16 wks		0-20 wks				
uon_		Male	Female	Average	Male	Female	Average	Male	Female	Average		
1	1/2SM+1/2Bj	0.89±0.07	0.54±0.08	0.71±0.05	29.21 <u>+</u> 0.49	23.04 ±0.59	26.42 ±0.39	35.94 ±5.72	29.23+4.60	33.16 <u>+</u> 6.23		
	BjBj	0.64+0.12	0.61±0.12	0.63±0.08	27.66 0.85	21.69 ±0.86	24.67 ±0.61	33.70 ±4.16	26.50±3.5	30.21 <u>+</u> 5.27		
Ov	erall mean	0.77 <u>+</u> 0.07	0.57+0.02	0.67+0.05	28.46+0.49	22.69 ±0.53	25.55±0.36 <sup>B</sup>	35.37 <u>+</u> 5.44	28.37+4. 45	32.33±6.11 <sup>B</sup>		
2	3/4SM+1/4Bj	0.63 <u>+</u> 0.06	0.61 <u>+</u> 0.06	0.62 <u>+</u> 0.04	27.64 <u>+</u> 0.42	22.93 ±0.44	25.28 ±0.31	37.70 ±5.95	30.09 <u>+</u> 5.35	33.75 <u>+</u> 6.65		
BjBj		0.59±0.10	0.59+0.09	0.59±0.06	27.01 ±0.70	22.38 ±0.64	24.70 ±0.47	33.74 ±4.05	28.13 <u>+</u> 3.63	30.63±4.73		
Ov	erall mean	0.61+0.06	0.60+0.05	0.60+0.03	27.32± 0.41	22.66 ±0.39	24.99 <u>+</u> 0.28 <sup>C</sup>	36.18±5.69	29.44 <u>+</u> 4.92	32.82 ±6.30 <sup>8</sup>		
3	7/8SM+1/8Bj	0.68+0.04	0.65±0.05	0.67±0.04	35.66 ±0.34	27.37 ±0.42	31.43 ±0.27	43.30±10.21	33.34+7.00	39.42 ±10.3		
	BjBj	0.60+0.19	0.61±0.18	0.60 <u>+</u> 0.13	26.87 ±1.45	22.80 ±1.27	24.80 ±0.96	32.81 ±5.11	27.86+3.39	30.08 <u>+</u> 4.87		
Ov	erall mean	0.64+0.10	0.62+0.09	0.63+0.07	31.18+ 0.74	25.09 ±0.67	28.14+0.50 <sup>A</sup>	42.76±10.20	32.82+6.91	38.77±10.29 <sup>A</sup>		
-	rall mean of raded groups	0.74 <u>+</u> 0.03	0.60 <u>+</u> 0.04	0.67±0.02	30.78 ±0.24	24.65 ±0.28	27.71 ±0.19ª	39.86 <u>+</u> 8.91	31.2 <del>9_6</del> .22	36.23 ±8.95*		
Overall	mean of Baheij breed	0.61 <u>+</u> 0.08	0.60 <u>+</u> .07	0.60 <u>+</u> 0.05	27.18 <u>+</u> 0.61	22.29 ±0.55	24.74 ±0.41 <sup>b</sup>	33.61 <u>+</u> 4.20	27.63 <u>+</u> 3.60	30.43 ±4.90b		
Ov	erall mean	0.67 <u>+</u> 0.04	0.60+0.04		28.98 <u>+</u> 0.33 <sup>X</sup>	23.47+0.31 <sup>V</sup>		38.78+8.62 <sup>X</sup>	30.29±5.90 <sup>V</sup>			
					Significanc							
Gen	eration (Gn)		NS			***			***			
Ger	Genotype (Gt) NS		NS			***		***				
	Sex (S)		NS		***			***				
Gn x Gt			NS			***		***				

SM: Silver Montazah strain and Bj: Baheij breed,

- Means having different letters in every column and within every factor of treatments are significantly different (P<0.05).

- The interactions between the Gn x S, Gt x S and Gn x Gt x S were not significant at the different periods.

Table (4): Effect of genotype and generation on least squares means and  $\pm$  standard error of body weight, some body measurements and body conformation of the two genotypes studied at 8 and 12 weeks of age for males and females of the 3<sup>rd</sup> generation

Age	Genotype	Sex	Body weight (g)	Shank length (cm)	Keel length (cm)	Breast width (cm)		
						`		
ļ	7/8SM+1/8Bj	M F	585.5 <u>+</u> 21.6	6.81±0.27	6.13 <u>+</u> 0.30	4.36+0.22		
		F	455.2±12.8			4.40 <u>+</u> 0.09		
	Average		494.5+13.3	6.83±0.13	6.11 <u>+0.14</u>	4.39±0.09		
	BiBi	M	499.0 <u>+</u> 20.2	7.44±0.32	6.76 <u>+</u> 0.41	5.00 <u>+</u> 0.24		
		F	445.1 <u>+</u> 10.2	6.98 <u>+</u> 0.14	6.2 <u>4+</u> 0.15	4.55 <u>+</u> 0.08		
8 wk	Average		452.4±9.5	7.04 <u>+</u> 0.13	6.31 <u>+</u> 0.14	4.61±0.08		
	Overall mean males		559.9 <u>+</u> 17.9	6.99 <u>+</u> 0.22	6.31 <u>+</u> 0.25	4.55 <u>+</u> 0.18		
	Overall mear females		449.8 <u>+</u> 8.0	6.91 <u>+</u> 0.10	6.18 <u>+</u> 0.11	4.48 <u>+</u> 0.06		
Over	all mean of 8 w	eeks	474.2 <u>+</u> 8.5 <sup>b</sup>	6.93 <u>+</u> 0.09 <sup>b</sup>	6.21±0.10 <sup>b</sup>	4.50 <u>+</u> 0.06		
		М	937. <u>4+</u> 29.2	7.97 <u>+</u> 0.19	7.58 <u>+</u> 0.22	4.77 <u>+</u> 0.15		
	7/8SM+1/8Bj	F	783.2 <u>+</u> 18.8	7.90 <u>+</u> 0.13	7.43±0.11	4.85±0.16		
	Average		832.1+18.2	7.92 <u>+</u> 0.10	7.48 <u>+</u> 0.10	4.82 <u>+</u> 0.08		
	D:D:	M	880.9 <u>+</u> 128.4	7.38 <u>+</u> 0.40	7.13±0.52	4.90±0.56		
	BjBj	F	726.4 <u>+</u> 65.3	7.08±0.17	6.75 <u>+</u> 0.25	4.93±0.34		
	Average		803.6+72.8	7.23 <u>+</u> 0.21	6.94 <u>+</u> 0.27	4.91 <u>+</u> 0.30		
12 wk	Overall mean of males		928.0 <u>+</u> 31.1	7.87 <u>+</u> 0.12	7.50 <u>+</u> 0.20	4.79 <u>+</u> 0.15		
• •	Overall mean of females		778.3 <u>+</u> 18.0	7.87 <u>+</u> 0.17	7.37 <u>+</u> 0.11	4.85±0.09		
Over	all mean of 12 w	eeks	828.9+17.9ª	7.84±0.10°	7.42+0.10 <sup>a</sup>	4.83+0.08		
-	10014-1001	М	766.0±33.8	7.40+0.19	6.87+0.22	4.57+0.13		
//	8SM+1/8Bj	F	617.2 <u>+</u> 21.0	7.36+0.11	6.67+0.12	4.62±0.07		
	Overall mean of 7/8SM+1/8Bj		663.3±18.8 <sup>A</sup>	7.37±0.10	6.79 <u>+</u> 0.10	4.61±0.06		
		М	626.3+68.0	7.42+0.24	6.88±0.31	4.97±0.23		
	BjBj	F	465.6 <u>+</u> 14.4	6.99+0.13	6.28+0.14	4.58±0.08		
Ov	erall mean of B	jBj	494.4+18.3 <sup>B</sup>	7.06+0.12	6.390+0.13	4.65+0.08		
	erall mean of Ma		733.1 <u>+</u> 31.2 <sup>X</sup>	7.40 <u>+</u> 0.15	6.87±0.18	4.66+0.12		
	rall mean of Fen		558.5±15.8 <sup>Y</sup>	7.22±0.09	6.57±0.09	4.60±0.05		
	Overall mean		604.7±15.0	7.27+0.08	6.65+0.08	4.62+0.05		
			Signi	ficance of::				
	Age		**	**	**	NS		
	Genotype		*	NS	NS	NS		
	Sex		**	NS	NS	NS		
	Age x Genotype		NS	**	*	NS		

Cont. Table (4)

				Daducant	Body conformation							
Age	Genotype	Sex				C4						
			C1	C2	C3							
	7/8SM+1/8Bj	M	1.27±0.06	89.10±5.31	1.43 <u>+0.08</u>	100.47 <u>+</u> 6.69						
	7705141-17053	F	1.14 <u>+</u> 0.02	67.20 <u>+</u> 1.88	1.29±0.03	75.89 <u>+</u> 2.38						
	Average		1.18 <u>+</u> 0.02	73.80 <u>+</u> 2.41	1.33+0.03	83.30 <u>+</u> 2.95						
8	BjBj	M	1.08+0.04	67.75±1.45	1.20±0.06	74.57 <u>+</u> 2.30						
wk	DjDj	F	1.11 <u>+</u> 0.02	64.06±1.14	1.25 <u>+</u> 0.03	72.19±1.54						
	Average	2	1.10+0.02	64.49 <u>+</u> 1.01	1.24+0.02	72.52 <u>+</u> 1.36						
	Overall mean	of males	1.21±0.05	82.63 <u>+</u> 4.21	1.36 <u>+</u> 0.06	92.80+5.25						
	Overall mean of	females	1.12±0.01	65.51±1.07	1.27 <u>+</u> 0.02	73.90±1.38						
Ov	erall mean of 8	weeks	1.14+0.02b	69.30±1.40 <sup>b</sup>	1.29 <u>+</u> 0.02	78.08±1.72 <sup>b</sup>						
	2/9Ch4+1/9D:	M	1.24+0.03	118.56+4.11	1.31±0.04	125.35 <u>+</u> 4.95						
	7/8SM+1/8Bj	F	1.18+0.02	100.18+2.87	1.25±0.02	106.25 <u>+</u> 2.93						
	Average	e	1.20+0.02	106.02+2.57	1.27+0.02	112.31+2.76						
1		M	1.31+0.14	123.51+24.94	1.37+0.02	129.14+26.55						
12	BjBj	F	1.27+0.06	103.20+10.95	1.34+0.16	107.83+9.30						
wk	Average	e	1.29+0.07	113.35+13.18	1.35+0.08	118.49+12.65						
	Overall mean of males		1.25 <u>+</u> 0.03	119.39±5.03	1.32 <u>+</u> 0.08	123.98 <u>+</u> 5.88						
)		Overall mean of										
1	females		1.18 <u>+</u> 0.02	100.44 <u>+</u> 2.75	1.26 <u>+</u> 0.02	106.38 <u>+</u> 2.77						
Ov	erall mean of 12		1.21+0.02*	106.84+2.69ª	1.28+0.02	113.01+2.85 a						
OV	CIAH MCAR OF 12	M	1.25+0.03	104.20+4.07	1.37+0.04	113.23+4.54						
	7/8SMx1/8Bj	F	1.16+0.02	83.50±2.46	1.27±0.02	90.89+2.49						
-	Overall mean		1.10±0.02	83.30 12.40	1.27±0.02	70.67 <u>1</u> 2.47						
	7/8SM+1/8E		1.19 <u>+</u> 0.02	89.91 <u>+</u> 2.27	1.30 <u>+</u> 0.02	97.81 <u>+</u> 2.39						
	BjxBj	M	1.15±0.06	86.00±11.02	1.26 <u>+</u> 0.07	92.77 <u>+</u> 11.25						
l	DixDi	F	1.12±0.02	66.90±1.88	1.26±0.03	74.78±1.99						
	Overall mean of	BjBj	1.13+0.02	70.33+2.61	1.26+0.02	78.00 <u>+</u> 2.68						
	Overall mean of	Males	1.23+0.03	99.93+4.13	1.34±0.04	108.41 <u>+</u> 4.48 <sup>X</sup>						
O	verall mean of F	emales	1.14+0.01	77.07±1.80	1.26±0.02	84.65±1.83 <sup>Y</sup>						
	Overall mea	n	1.17+0.01	83.11+1.86	1.28+0.01	90.93 <u>+</u> 1.94						
			Signif	icance of::								
	Age		**	**	NS	**						
	Genotype		NS	NS	NS	NS						
	Sex		NS	**	NS	**						
	Age x Genoty	pe	*			*						
_												

SM: Silver Montazah strain, Bj: Baheij breed, M: Male, F: Female,

<sup>-</sup> Means having different letters in every column and within every main factor of treatments are significantly different (P<0.05).

<sup>-</sup> Genotype had no significant effect on all traits studied except that for body weight where it affected significantly (P<0.01), and the interactions of Age x Genotype, Age x Sex, and Genotype x Sex had no significant effects on all traits studied.

Table (5): Effect of genotype and generation on least squares means and ± standard error of viability of male and female birds at different periods studied

				<u> </u>						
l				Chicks viability						
Gene-	Genetyne	0-4 wks		4-8 wks						
ration	. Genotype	Non sexed	Males	Females	Average	Non sexed				
1	1/2SM+1/2Bj	89.42 <u>+</u> 2.39	98.89 <u>+</u> 1.65	100.00 <u>+</u> 1.65	99.44 <u>+</u> 1.16	88.91 <u>+</u> 2.32				
1	BjBj	95.56 <u>+</u> 5.35	100.00±3.69	100.00±3.20	100.00±2.44	97.15±5.20				
Ove	erali mean	92.49 <u>+</u> 2.93	99.44+2.02	100.00 <u>+</u> 1.08	99.72 <u>+</u> 1.35 <sup>A</sup>	93.03 <u>+</u> 2.85				
2	3/4SM+1/4Bj	94.07 <u>+</u> 2.32	95.28 <u>+</u> 1.60	97.32±1.60	96.30 <u>+</u> 1.31	90.30 <u>+</u> 2.25				
2	BjBj	95.97 <u>+</u> 4.63	98.21 <u>+</u> 3.20	95.39±3.20	96.80 <u>+</u> 2.26	92.98+4.51				
Ove	rall mean	95.02 <u>+</u> 2.59	96.75 <u>+</u> 1.78	96.36±1.78	96.52 <u>+</u> 1.26 <sup>B</sup>	91.59+2.52				
2	7/8SM+1/8Bj	94.22 <u>+</u> 2.39	100.00±1.65	98.84±1.65	98.84±1.65 99.42±1.17					
3	BjBj	97.92 <u>+</u> 4.63	100.00±3.20	100.00±3.69	100.00±2.44	95.99±4.51				
Ove	rall mean	96.07 <u>+</u> 2.61	100.00+1.80	99.42+2.02	99.71 <u>+</u> 1.35 <sup>A</sup>	94.92+2.54				
	all mean of ided groups	92.57 <u>+</u> 1.37	98.06 <u>+</u> 0.94	98.72 <u>+</u> 0.94	98.39 <u>+</u> 0.66	90.98±1.33				
	mean of Baheij breed	96.48 <u>+</u> 2.82	99.40 <u>+</u> 1.94	99.40±1.94 98.46±1.94		95.73 <u>+</u> 2.74				
Ove	rali mean	96.07 <u>+</u> 2.61	98.73±1.08	98.59±1.08	98.44	94.92+2.54				

SM: Silver Montazah strain and Bj: Baheij breed,

- Means having different letters in every column and within every factor of treatments are significantly different (P<0.05),

- All main factors studied and all interactions between them were not significant except generation which affected (P<0.01) viability at 4-8 period.

Table (6): Effect of genotype and generation on least squares means and ± standard error of live body weight, g and initial weights (g) and percentages of carcass traits at 16 and 20 weeks age at the 2<sup>nd</sup> and 3<sup>rd</sup> generations

		A ==	Line					Carcas	traits		
Generation	Genotype	Age, wks	Live b			Caro				Le	gs
		wks	weig	nt j	Wt		%		W	_	%
	3/4SM+1/4Bj	. 16	1014+	13	663±1	8	65.4	<u>+</u> 1.1	66.9	±3.1	6.59±0.29
	3/43 WIT 1/4BJ	20	1336+	15	951±3	0.	71.1	<u>+</u> 1.3	74.4	±3.4	5.58 <u>+</u> 0.27
	Average		1175+	10	794 <u>+</u> 4	8	68.0		70.7		6.08±0.18
	BjBj	16	945+	14	611 <u>+</u> 1	7	64.7		58.6	±3.4	6.21±0.27
2	5,5,	20	1181	:15	816 <u>+</u> 1	7	69.1	<u>+</u> 1.1	63.7	<u>+</u> 3.2	5.40 <u>+</u> 0.30
	Average		10634	11	714+3	6	66.9	<u>+</u> 1.3	61.2	+2.2	5.80±0.19
	Overall mean a		980 <u>+</u>	10	640 <u>+</u> 1	4	65.1	65.1 <u>+</u> 1.0		<u>+</u> 2.3	6.40 <u>+</u> 0.18
	Overall mean at 20 wk		1259	12	884 <u>+</u> 2	27	70.1	0.1 <u>+</u> 0.9		<u>+</u> 2.4	5.49 <u>+</u> 0.19
Overall mean of generation 2			1119	7 b	756 <u>+</u> 3	1 <sup>b</sup>	67.5	±0.8	65.9	<u>+</u> 1.7	5.94±0.13
	7/8SM+1/8Bj	16	1348-	15	912±3	33	67.6	±1.8	59.4	<u>+</u> 3.4	4.41±0.27
	//65WITI/6BJ	20	1602-	12	1150±	28	71.8	±1.7	77.7	±2.9	4.85±0.23
	Average		1475	10	1051 <u>+</u> 40		70.0	±1.3	68.6	<u>+2.2</u>	4.63+0.17
3	BjBj	16	974 <u>+</u> 15		644 <u>+</u>	11	66.1	±1.1	54.7	<u>+</u> 3.4	5.62±0.27
3		20	1186 <u>+</u> 17		825±2	23	69.6	±04	76.2	±3.6	6.42 <u>+</u> 0.30
	Average		1080			30	67.8	<u>+0.8</u>	65.4	+2.4	6.02+0.19
	Overall mean at 16 wk		1161;	161±10 77		47 66.9		±1.0 57.0		<u>+2.4</u>	5.01±0.19
	Overall mean at	20 wk	1394	l±12 1014±					77.0 <u>+</u> 2.2		5.64 <u>+</u> 0.18
Overall n	nean of generati	on 3	1278		907+42*		69.1+0.8				5.32+0.13
Overall m	ean of 16 wks o	f age	1070	<u>+</u> 7 <sup>¥</sup>	706 <u>+</u> 2	8 <sup>Y</sup>	65.9+0.7 <sup>X</sup>		59.9+1.7 Y		5.70±0.11
Overall m	ean of 20 wks o	f age	1328+		955 <u>+</u> 3		70.5	±0.6 <sup>Y</sup>	73.0	<u>+1.6 <sup>X</sup></u>	5.56±0.13
Overall me	an of upgraded	group			928 <u>+</u> 4	1^	69.1	±0.9	70.0	<u>+1.6 ^</u>	5.4±0.2 <sup>B</sup>
Overall m	ean of Baheij l	oreed	1072	<u>+</u> 7₿	724+2	3 <sup>B</sup>	67.4	±0.7	63.3	±1.7 <sup>8</sup>	5.9 <u>+</u> 0.1 <sup>A</sup>
0	verall mean		121		833		68	3.3	6	7.0	5.6
				Sign	ificance						
	Generation (G	n)			**	,	**	Z	S	NS	**
	Genotype (Gt	()			**		**	Z		*	**
Age					**		**	**	*	**	NS
	Gn x Get				**	-	**	NS		NS	**
	Gn x Age				*	_	NS		NS		**
Gt x Age					**		**	Z	S	NS	NS
	Gn x Gt x Ag	e			NS	ı	NS	N	S	NS	NS

SM: Silver Montazah strain, Bj: Baheij breed, M: Male, F: Female,

<sup>-</sup> Means having different letters in every column and within every main factor of treatments are significantly different (P<0.05).

Cont. Table (6):

j., -

Genotype attion   Genotype   Age, wks   Gizzard   Liver   Heart						Carcass	traits		
3/4SM+1/4Bj	Gener-	Genotype		Gizz	ard	Liv		He	art
Average	ation	,	WKS		%				
Average		2/45N4+1/4D:	16 .	38.5±2.4	3.8 <u>+</u> 0.2	28.2 <u>+</u> 1.5	2.8 <u>+</u> 0.1	8.8 <u>+</u> 0.9	0.87 <u>+</u> 0.1
BjBj		3/45M±1/4BJ	20	43.1 <u>+</u> 2.7	3.2±0.2	30.8±1.6	2.3 <u>+</u> 0.1	10.1 <u>+</u> 0.9	0.76 <u>+</u> 0.1
Average		Averag	ge	40.8 <u>+</u> 1.8	3.5±0.2	29.5±1.1	2.5±0.1	9.4 <u>+</u> 0.6	0.81 <u>+</u> 0.1
20   31.5±2.7   2.7±0.2   33.2±1.8   2.8±0.1   10.4±0.9   0.82±0.1		D:D:	16	30.6 <u>+</u> 2.7	3.2±0.2	24.6 <u>+</u> 1.6	2.6±0.1	7.04 <u>+</u> 0.9	0.75 <u>+</u> 0.1
Overall mean at 16 wk         34.6±1.8         3.5±0.2         26.4±1.1         2.7±0.1         7.9±0.6         0.81±0.1           Overall mean of generation 2         35.9±1.3         3.2±0.8         29.2±0.8         3.2±0.1         9.1±0.4         0.82±0.           Overall mean of generation 2         35.9±1.3         3.2±0.8         29.2±0.8         3.2±0.1         9.1±0.4         0.81±0.1           Average         44.1±2.7         3.3±0.2         27.0±1.6         2.0±0.1         9.6±0.9         0.71±0.1           Average         43.0±1.8         2.5±0.2         26.6±1.4         1.7±0.1         15.9±0.8         1.0±0.1           BjBj         16         32.0±2.7         3.3±0.2         22.1±1.6         2.3±0.1         7.1±0.9         0.73±0.           Average         37.2±1.9         3.4±1.2         29.1±1.1         2.7±0.1         9.7±0.6         0.88±0.           Overall mean at 16 wk         38.1±1.9         2.3±0.2         24.6±1.1         2.1±0.1         8.3±0.6         0.72±0.           Overall mean of generation 3         40.1±1.3         3.2±0.2         24.6±1.1         2.1±0.1         8.3±0.6         0.72±0.           Overall mean of lowks of ag		ВЈВЈ	20	31.5±2.7	2.7_0.2	33.2 <u>+</u> 1.6	2.8±0.1	10.4 <u>+</u> 0.9	0.88±0.1
Overall mean at 20 wk   37.3±1.9   3.0±0.1   32.0±1.1   2.6±0.1   10.2±0.6   0.82±0.	2	Avera	ge	31.2 <u>+</u> 1.9	3.0±0.2	28.9 <u>+</u> 1.1	2.7 <u>+</u> 0.1	8.7 <u>+</u> 0.6	$0.82 \pm 0.1$
Nerall mean of generation 2   35.9±1.3   3.2±0.8   29.2±0.8   3.2±0.1   9.1±0.4*   0.81±0.1		Overall mean	at 16 wk	34.6 <u>+</u> 1.8	3.5±0.2	26.4±1.1	2.7 <u>+</u> 0.1	7.9 <u>+</u> 0.6	0.81 <u>+</u> 0.1
T/8SM+1/8Bj   16					3.0 <u>+</u> 0.1	32.0 <u>+</u> 1.1	2.6±0.1		0.82 <u>+</u> 0.1
T/8SM+1/8Bj   16	Overal				3.2 <u>+</u> 0.8	29.2±0.8	3.2±0.1	9.1±0.4 <sup>b</sup>	0.81 <u>+</u> 0.0
Average					3.3 <u>+</u> 0.2	27.0±1.6	2.0±0.1		0.71±0.1
BjBj		//85M+1/8BJ	20	42.0 <u>+</u> 2.3	2.6±0.2	26.6 <u>+</u> 1.4	1.7±0.1	15.9 <u>+</u> 0.8	1.0 <u>+</u> 0.1
Average	Ì	Avera	ge	43.0+1.8	2.5±0.2	26.8 <u>+</u> 1.1	1.8 <u>+</u> 0.1	12.7 <u>+</u> 0.6	0.85±0.1
Average		BjBj	16	32.0 <u>+</u> 2.7	3.3±0.2	22.1±1.6	2.3±0.1	7.1±0.9	0.73 <u>+</u> 0.1
Overall mean at 16 wk   38.1±1.9   2.3±0.2   24.6±1.1   2.1±0.1   8.3±0.6   0.72±0.			20	42.4 <u>+</u> 2.7	3.6±0.2	36.0±1.6	3.0 <u>+</u> 0.1	12.2 <u>+</u> 0.9	1.03 <u>+</u> 0.1
Overall mean at 16 wk   38.1±1.9   2.3±0.2   24.6±1.1   2.1±0.1   8.3±0.6   0.72±0.	1 2	Avera	ge	37.2±1.9	3.4 <u>+</u> 1.2	29.1±1.1	2.7 <u>+</u> 0.1	9.7 <u>+</u> 0.6	0.88 <u>+</u> 0.1
Overall mean of generation 3         40.1±1.3         3.2±0.8         27.9±0.8         3.2±0.1         11.0±0.4*         0.87±0.0           Overall mean of 16 wks of age         36.3±1.3         3.4±0.1         25.5±0.8*         2.4±0.1         8.1±0.4*         0.76±0.0           Overall mean of 20 wks of age         39.7±1.3         3.0±0.1         31.7±0.8*         2.5±0.1         12.1±0.4*         0.92±0.0           Overall mean of upgraded groups         41.9±1.3 x         3.2±0.1         28.1±0.75         2.2±0.1         11.1±0.4 x         0.83±0.0           Overall mean of Baheij breed         34.1±1.3 x         3.2±0.1         29.1±0.8         2.7±0.1         9.2±0.5 x         0.84±0.0           Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         NS           Age         NS         NS         NS         NS         NS           Age         NS         **         NS         NS         NS           Gn x Age         NS         NS         NS         NS         NS           Gt x Age         NS         NS         NS	3	Overall mean at 16 wk		38.1 <u>+</u> 1.9	2.3 <u>+</u> 0.2	24.6 <u>+</u> 1.1	2.1 <u>+</u> 0.1	8.3±0.6	0.72 <u>+</u> 0.1
Overall mean of 16 wks of age         36.3+1.3         3.4+0.1         X         25.5+0.8         2.4+0.1         8.1+0.4         0.76+0.0           Overall mean of 20 wks of age         39.7+1.3         3.0+0.1         11.7+0.8         2.5+0.1         12.1+0.4         0.92+0.0           Overall mean of upgraded groups         41.9+1.3         3.2+0.1         28.1+0.75         2.2+0.1         11.1+0.4         0.83+0.0           Overall mean of Baheij breed         34.1+1.3         3.2+0.1         29.1+0.8         2.7+0.1         9.2+0.5         0.84+0.0           Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         NS           Age         NS         **         NS         NS         **         NS           Age         NS         **         NS         NS         NS         NS           Gn x Age         NS         NS         NS         NS         NS         NS           Gt x Age         NS         NS         NS         NS         NS         NS         NS		Overall mean	at 20 wk	42.2 <u>+</u> 1.8	3.1±0.1	31.3±1.1	2.4±0.1	14.1±0.6	1.01 <u>+</u> 0.1
Overall mean of 20 wks of age         39.7±1.3         3.0±0.1         31.7±0.8         2.5±0.1         12.1±0.4         0.92±0.0           Overall mean of upgraded groups         41.9±1.3         3.2±0.1         28.1±0.75         2.2±0.1         11.1±0.4         0.83±0.0           Overall mean of Baheij breed         34.1±1.3         3.2±0.1         29.1±0.8         2.7±0.1         9.2±0.5         0.84±0.0           Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         NS           Genotype (Gt)         **         NS         NS         NS         NS           Age         NS         **         **         **         **           Gn x Gt         NS         **         NS         NS         NS           Gr x Age         NS         NS         NS         NS         NS	Overa	ll mean of gene	ration 3			27.9±0.8	3.2±0.1	11.0+0.4*	0.87 <u>+</u> 0.0
Overall mean of 20 wks of age         39.7±1.3         3.0±0.1         31.7±0.8         2.5±0.1         12.1±0.4         0.92±0.0           Overall mean of upgraded groups         41.9±1.3         3.2±0.1         28.1±0.75         2.2±0.1         11.1±0.4         0.83±0.0           Overall mean of Baheij breed         34.1±1.3         3.2±0.1         29.1±0.8         2.7±0.1         9.2±0.5         0.84±0.0           Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         NS           Genotype (Gt)         **         NS         NS         NS         NS           Age         NS         **         **         **         **           Gn x Gt         NS         **         NS         NS         NS           Gr x Age         NS         NS         NS         NS         NS	Overa	ll mean of 16 w	ks of age	36.3+1.3	3.4 <u>+</u> 0.1 <sup>X</sup>			8.1 <u>+</u> 0.4	0.76 <u>+</u> 0.0
Overall mean of upgraded groups         41.9±1.3 A         3.2±0.1         28.1±0.75         2.2±0.1         11.1±0.4 A         0.83±0.0           Overall mean of Baheij breed         34.1±1.3 B         3.2±0.1         29.1±0.8         2.7±0.1         9.2±0.5 B         0.84±0.0           Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         NS           Genotype (Gt)         **         NS         NS         NS         NS           Age         NS         **         **         *         **         **           Gn x Gt         NS         **         NS         NS         NS         NS           Gt x Age         NS         NS         NS         NS         NS         NS	Overs	ll mean of 20 w	ks of age	39.7±1.3	3.0±0.1 Y	31.7±0.8 X	2.5±0.1 X	12.1 <u>+</u> 0.4	0.92 <u>+</u> 0.0 <sup>X</sup>
Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         **         NS           Genotype (Gt)         **         NS         NS         **         NS         NS         **         NS           Age         NS         **         **         **         **         **         **         **         **           Gn x Gt         NS         **         NS         **         NS	Overal	mean of upgrade	ed groups	41.9±1.3 A	3.2 <u>+</u> 0.1	28.1±0.75	2.2±0.1		0.83 <u>+</u> 0.0
Overall mean         38.2         3.2         28.5         2.4         10.4         0.85           Significance of:           Generation (Gn)         *         NS         NS         NS         **         NS           Genotype (Gt)         **         NS         NS         **         NS         NS         **         NS           Age         NS         **         **         *         **         **         **         **         **           Gn x Gt         NS         **         NS         **         NS	Overa	all mean of Bahe	ij breed	34.1±1.3 <sup>B</sup>	3.2±0.1	29.1±0.8	2.7±0.1	9.2±0.5 <sup>B</sup>	0.84+0.0
Significance of:           Generation (Gn)         *         NS         NS         NS         **         NS           Genotype (Gt)         **         NS         NS         **         NS         **         NS           Age         NS         **         **         *         **         **         **           Gn x Gt         NS         **         NS         **         NS         NS         NS           Gn x Age         NS         NS         NS         NS         NS         NS         NS		Overall mean	n			28.5	2.4	10.4	0.85
Genotype (Gt)         **         NS         NS         **         NS           Age         NS         **         **         **         **         **         **         **         **         **         **         **         **         NS				Sig	gnificance	of:			
Genotype (Gt)         **         NS         NS         **         NS           Age         NS         **         **         **         **         **         **         **         **         **         **         **         NS		Generation (G	n)	*	NS	NS	NS	**	NS
Age         NS         **         **         **         **         **         **         **         **         **         **         **         MS         NS		Genotype (G	:)	**	NS		NS	**	
Gn x Gt         NS         **         NS         **         NS         NS           Gn x Age         NS         *         **         NS         **         **           Gt x Age         NS         NS         NS         NS         NS         NS					**	**		**	**
Gt x Age NS NS NS NS NS NS					**		**		
Gt x Age NS NS NS NS NS NS					*	**	NS	**	**
					NS	NS		NS	NS
			e e	*	*	NS	NS	NS	NS

SM: Silver Montazah strain, Bj: Baheij breed, M: Male, F: Female,

<sup>-</sup> Means having different letters in every column and within every main factor of treatments are significantly different.

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# الملخص العربى تحسين الصفات الإنتاجية والتكاثرية لدجاج بهيج بواسطة الخلط تأثير التزاوج القمى على:

أ. صفات النمو والحيوية وتناسق أعضاء الجسم وصفات الذبيحة

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تم إجراء دراسة لتقييم تأثير الخلط الرجعي (زيادة دم سلالة المنتزه الفضي) على بعض الصفات الإنتاجية (وزن الجسم – معدل النمو – كفاءة النمو)] وكذلك نسبة حيوية الطيور عند أعمار مختلفة ، وذلك من خلال خلط ديوك المنتزه الفضى المنتخبة لوزن الجسم الناضج كخط أباء مع إناث البهيج لإنتاج ثلاث أجيال متتابعة من الأبناء كما يلى: الجيل الأول (٢/١ منتزة فضي + ٢/١ بهيج) ، الجيل الثالث (٨/٧ مبتزة فضي + ٢/١ بهيج) ، الجيل الثالث (٨/٧ مبتزة فضي + ٢/١ أسبوع وأيضا صفات النبيحة عند ١٢/١ مبوع من العمر، وكانت النتائج كما يلى: -

- ١- تغوقت طيور مجموعة الخلط الرجعى في وزن الجسم معنويا عن طيور المجموعة المقارنة (كنترول) عند جميع الأعمار المدروسة وكذلك كانت طيور الجيل الثالث الأعلى معنويا في وزن الجسم مقارنة بطيور الجيلين الأول والثاني.
- ٢- أدى استخدام الخلط الرجعى إلى تحسين نسبة معدل النمو من الفقس حتى ١٦ أسبوع من عمر الطيور بينما وجد العكس خلال الفترة ٨-١٢ أسبوع من العمر. حقق الجيل الثالث أعلى نسبة معدل نمو خلال الفترات من الفقس إلى ١٦ أسبوع ، وأيضا من ١٦-١٦ أسبوع من العمر.
- ٣- تفوقت طيور مجموعة الخلط الرجعي في كفاءة النمو على طيور مجموعة بهيج خلال فترات النمو من الفقس إلى أربعة أسابيع ، ١٢ ، ١٦ أسبوع من العمر. كانت طيور الجيل الثالث الأعلى معنويا في كفاءة النمو خلال الفترات ٤-٨ ، من الفقس إلى ١٦ أسبوع ، ١٦ . ١٦ ، من الفقس إلى ١٦ أسبوع وكذلك من ١٦-٢٠ أسبوع من عمر الطيور. وكانت متوسطات الذكور أعلى معنويا عن متوسطات الإناث بالنسبة لصفات النمو المدروسة.
  - ٤- تأثرت الحيوية معنويا بالجيل خلال الفترة ٤-٨ أسبوع من العمر.
- حان تأثیر العمر معنویا علی کل من وزن الجسم ، طول الساق وأیضا طول عظمة القص وأیضا تفوقت طیور الجیل الثالث عن طیور الکنترول ، من ناحیة أخری لم تتأثر أی من صفات تکوین الجسم بالخلط الرجعی.
- ٦- تفوقت طيور الجيل الثالث على طيور الجيل الثانى بالنسبة لوزن الذبيحة، والأعضاء الداخلية الماكولة ، وأيضا تفوقت مجموعة الخلط الرجعى بالنسبة لجميع الصفات المدروسة مقارنة بطيور سلالة بهيج. أيضا كانت الاختلافات معنوية بين متوسطات الصفات المدروسة عند ١٦ ، ٢٠ أسبوع من العمر.