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Breeding for the improvement of indigenous chickens in Bangladesh: performance of second generation

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Abstract: The improvement of productivity of indigenous chicken is a long desire in the country. Present research is a part of the long-term selection program being undertaken to evaluate the carcass characteristics and expected response to selection of second generation (G_2) of indigenous chicken under intensive management in Bangladesh. A total of 1643-day-old chicks comprising of 3 types of chicken namely Naked Neck (NN), Hilly (H) and Non-descript Desi (ND) were hatched in a two batches for this study. In second generation (G_2), selection was practiced at 3 (three) stages of birds life, firstly and secondly at 8 and at 16 weeks of age, selection was performed on the basis of breeding value for 8 and 16 week body weight. Thirdly, at 40-week of age, on the basis of an index comprising the parameters of age at sexual maturity (ASM), body weight (BW), egg production (EP) and egg weight (EW). Improvement target of egg weight was to increase by 1g and improvement target of egg production rate was to increase by 2 % per generation. The main target was the mean body weight in H, ND and NN chicken has to be gone up from the initial eight-week body weight of 375, 342 and 331g to 500 g at eight weeks of age. At 8, 10 and 12 weeks of age, six birds from each genotype were slaughtered to analyze the meat yield traits. Slaughter data were analyzed in a 3 (genotype) \times 3 (slaughter age) factorial arrangement in CRD by General Linear Model (GLM) Univariate Procedure in SPSS Computer Program. Genotype and slaughter age had significant effect ($p < 0.001$) on dressing percentage. Body weight at 8 weeks of age was expected to improve by 51.21 vs. 24.03; 37.74 vs. 15.47 and 26.26 vs. 9.65g; respectively for ND, H and NN males and females. In terms of body weight H genotype was superior and NN genotype was for dressing percentage. As a result of selection; EP, BW increased and ASM reduced in second generation than that of the foundation stock.

Keywords: indigenous chicken; generation; selective breeding; breeding value; performance

1. Introduction

The improvement of productivity of indigenous chicken is a long desire in the country. Indigenous chickens of Bangladesh are categorized as Non-descriptive Deshi (ND), Naked Neck (NN), Hilly (H), Aseel (AS) and Jungle fowl (Bhuiyan *et al.*, 2005) in respect of the morphological variations as well as production performances. Bangladesh Livestock Research Institute (BLRI) since its inception did initiate programs for the conservation and development of indigenous chicken through several poultry development projects. As a part of

selection and improvement of indigenous chicken, this work in the form of a Ph.D. research was being planned. It is a long-term vision. As a first step, foundation stock was established utilizing the existing stock of Bangladesh Livestock Research Institute (BLRI) as well as by incorporating variation through screening of males/females/eggs from a wider indigenous chicken gene pool of Bangladesh. Response observed in most of the selection experiments with egg number as a selection criterion suggested a negative genetic correlation between egg number and egg weight (Hazary *et al.*, 1990). Mature body weight is closely related to initial egg weight, with heavier pullets producing heavier eggs (Prasad, 2000). Though early maturity and high rate of production are desirable goals, the gain in response may be at the expense of persistency if long enough records are not included in the selection. However, the present study was undertaken to evaluate the performance and expected response to selection of second generation (G_2).

2. Materials and Methods

2.1. Production of second generation (G_2)

The study was conducted at Bangladesh Livestock Research Institute, under Poultry Production Research Division, Savar, Dhaka, Bangladesh. A total of 1643 progenies (ND=926, H= 383 and NN= 334) were hatched in a two batches for second generation (G_2). Day-old chicks collected from the hatchery were weighed and leg banded individually.

2.2. Rearing and data collection of second generation (G_2)

All the chicks of three genotypes were transferred into the brooder, which were cleaned and disinfected earlier and 5% glucose solution was supplied for the first three days. One-week later leg bands were removed and wing bands were provided all the experimental birds. Debeaking was performed after 10-12 days of age. Concentrate mixtures that contain 20.06% Crude Protein & 2908 Kcal ME/kg DM; 18.13% Crude Protein & 2904 Kcal ME/kg DM and 16.33% Crude Protein & 2845 Kcal ME/kg DM were provided twice daily in the morning and evening during brooding, growing and laying period, respectively. Cool clean drinking water was supplied all the times. Water also was provided *ad libitum* twice daily in the morning and evening. Feeder and drinker were cleaned twice in a week. Refusals of the feed were measured everyday in the morning. All chicks were vaccinated as per schedule given by veterinarian. All productive and reproductive records are being kept.

2.3. Selection and mating plan of second generation (G_2)

The selection objectives of the study were to improve the egg production, egg weight and / or growth rate of indigenous chicken depending on the genotype of birds. Improvement target of egg weight is to increase by 1 g and improvement target of egg production rate is to increase by 2% per generation. The main target was the mean body weight in ND, H, NN chickens have to be gone up from the initial eight-week body weight of 342, 375, 331 g on an average 349g to 500 g at 8 week of age after 3 generations of selective breeding. In second generation (G_2), selection was practiced at 3 (three) stages of birds life, firstly and secondly, at 8 and at 16 weeks of age, selection was performed on the basis of breeding value for 8 and 16 week body weight. At 8 weeks of age, a total of 460 females (ND=230, H= 115 and NN= 115) and 175 males (ND=85, H= 45 and NN= 45) were selected according to breeding value on the basis of 8 week's body weight. At 16 weeks of age, a total of 400 females (ND=200, H= 100 and NN= 100) and 80 males (ND=40, H= 20 and NN= 20) were selected according to breeding value on the basis of 16 week's body weight. Thirdly, at 40-week of age, selection was done using multi trait index selection and independent culling level. The selection Index was computed by the following equation:

$$\text{Selection Index (I)} = b_1x_1 + b_2x_2 + \dots + b_nx_n$$

Where, x_1, x_2, \dots, x_n represent the phenotypic value for the trait

b_1, b_2, \dots, b_n denote the relative weight given to each of the trait

The total score was obtained from above calculation is a selection index. The individual with the higher total score was selected for breeding purposes. Both index selection and independent culling levels were used to select chicken. Culling level used was as follows:

Trait	Culling level
ASM	$X \leq AV+2SD$
BW	$AV-2SD \leq X \leq AV+2SD$
EW	$AV-SD \leq X \leq AV+SD$
EP	$AV \leq X$
Index	$AV \leq X$

A total of 37 males (ND=17, H= 10 and NN= 10) and 185 females (ND=85, H= 50 and NN= 50) were selected comprising the parameters of Age at first egg (days), Body weight (g) at Egg production percentage (168-280 days), 40 weeks of age and Egg Weight (g) at 38 - 40 weeks of age

A total of 20 males (ND=10, H= 5 and NN= 5) and 40 female birds (ND=20, H= 10 and NN= 10) were kept as spare birds. An assortative mating design was followed to keep inbreeding as low as possible.

2.4. Breeding design and experimental chicken

Name of genotype	Sex	No. of day old chicks	No. of growing chicks		No. of adult birds	No. of selected bird at 40 wks of age	
			8 wks	16 wks.		Selected	Spare
ND	Male	926	85	40	40	17	10
	Female		230	200		85	20
H	Male	383	45	20	20	10	5
	Female		115	100		50	10
NN	Male	334	45	20	20	10	5
	Female		115	100		100	50

2.5. AI and collection of hatching eggs to produce third generation (G₃)

Selected males and females were mated using artificial insemination. Artificial Insemination was practiced twice in a week. Hatching eggs were collected up to 10 days. Male birds were collected from different parts of Bangladesh to introduce new variability. Two batches were required to produce third generation (G₃).

2.6. Data recording

All productive and reproductive parameters were recorded for second generation (G₂). Records were kept on day-old weight (g), fortnightly individual body weight up to 8 weeks, monthly weight up to 20 weeks, daily egg production, and egg weight at 38-40 weeks of age, temperature and humidity, growth rate, feed intake and feed conversion ratio (FCR). Feed conversion ratio (FCR) was recorded for the whole period as total feed intake (kg) per kg weight gain. Temperature and humidity were recorded four times a day; (06:00h, 12:00h, 18:00h and 24:00h).

2.7. Slaughtering and carcass characteristics data

At 8, 10 and 12 weeks of age, six males of each genotype were slaughtered to analyze the meat yield traits. All chickens were kept off feed overnight before slaughtering but drinking water was provided ad libitum. Birds were slaughtered following 'halal' method (Singh *et al.*, 2003) by severing the jugular vein allowed to bleed completely and then plucked and weighed to determine blood and feather losses (Kotula *et al.* 1960; Pandey and Shyamsunder, 1990). Pre-slaughter live weight, blood loss weight, eviscerated weight, breast meat weight, thigh plus drumstick weight etc. were recorded. All weight related to carcass characteristics were expressed as the percentage of live weight. Carcasses were dissected according to Singh *et al.* (2003) except that birds were not scalded.

2.8. Statistical analysis

The treatments were arranged in a 3 (genotype) × 3 (slaughter age) factorial experiment. The data were analyzed by factorial arrangement in a CRD by General linear Model (GLM) Univariate Procedure in SPSS Computer Program. The following general linear statistical models were used to analyze the different parameters:

i) $Y_{ij} = \mu + g_i + e_{ij}$, where, Y_{ij} is the dependent variable of the experiment; μ is the overall mean; g_i is the effect of i th genotype ($i=1-3$); e_{ij} is the error term specific to each record.

ii) $Y_{ij} = \mu + s_i + e_{ij}$, where, Y_{ij} is the dependent variable of the experiment; μ is the overall mean; s_i is the effect of i th slaughter age ($i=1-3$); e_{ij} is the error term specific to each record.

2.9. Prediction of expected selection response

Expected selection response in three types of indigenous chicken for body weight at 8 weeks was estimated using the following equation (Falconer, 1981).

$$i) R = h^2 \times S$$

Where,

R = Expected response in mass selection

h^2 = heritability, h^2 for BW at 8 weeks of age

S = Selection differential

3. Results and Discussion

3.1. Carcass characteristics

Dressing percentage ($p < 0.001$) and live weight ($p < 0.05$) was affected by genotype (Table 1). Significantly ($P < 0.001$) higher dressing percentage was found in NN (67.29) genotype than that of ND (66.92) and H (65.4) genotypes. Similar result was found in NN (64.58) genotype than that of ND (60.26) and H (61.70) genotypes (Faruque *et al.*, 2011). Iqbal *et al.* (2009) found the average live weight 1.72 ± 0.04 kg and 1.25 ± 0.02 kg for cock and hen respectively, which differed significantly ($p < 0.01$) between the two sexes. They also reported that dressing percentage of 70.11 ± 0.66 for cock and 63.80 ± 1.59 for hen were significantly different ($p < 0.01$). There was significant ($P < 0.001$) difference in live weight and dressing percentage between ages at slaughter (Table 2).

3.2. Selection differential, selection intensity and expected response to selection

Selection differential at 8 weeks of age for ND, H and NN males were 102.63, 77.20 and 54.94 g respectively (Table 3). The corresponding values for females were 48.18, 31.65 and 20.19 g. The intensity of selection for males at 8 weeks of age was 4.73, .84 and 0.96 for ND, H and NN genotypes. In females, the corresponding figures were 4.35, 1.64 and 0.43. Genotype wise expected response to selection for 8 week body weight is shown in Table 3. As a result of selection, body weight at 8 weeks of age was expected to improve by 51.21 vs. 24.03; 37.74 vs. 15.47 and 26.26 vs. 9.65g; respectively for ND, H and NN males and females. Depending on a selective breeding, results from the present study indicate that on an average weight gain is 27 g and it would take approximately 5 to 6 generations of selection, which corresponds to around 3 to 4 years of selection for improving body weight at 8 week from their current population mean of 349 g to the desired population mean of 500 g.

3.3. Comparison between second generation (G_2) and foundation stock (G_0)

Weight gains in indigenous chicken at hatch and at 8 week of age for G_2 , were -1.97, -1.94, 0.44 g and 107.34, 175.95, 150.70 g; respectively for ND, H and NN genotypes (Table 4). It was observed that all genotypes came to sexual maturity as 8 to 19 days earlier in the second generation (G_2) than that of the foundation stock (G_0). Egg production increased as 3 to 14 numbers in the second generation (G_2) than that of the foundation stock (G_0). Egg weight decreased in ND (-0.48 g) and NN (-0.324 g) but increased in H (1.47 g). Body weight increased by 202.91, 337.36 and 72.82 g at 40 week of age; respectively for ND, H and NN genotypes. Singh and Singh (1989) reported that as a result of selection egg production is expected to improve by 6.76 eggs with minimum decrease in age at first egg (1.46 days) and marginal decline in egg weight (0.127 g) per generation.

Table 1. Effect of genotype on carcass characteristics.

Parameter	Genotype			P-value
	ND Mean \pm SD	H Mean \pm SD	NN Mean \pm SD	
Live weight (g)	756.50 ^b \pm 157.62	798.50 ^a \pm 198.38	763.16 ^b \pm 174.72	$P < 0.05$
Dressing %	66.92 ^a \pm 1.77	65.40 ^b \pm 2.50	67.29 ^a \pm 1.79	$P < 0.001$

^{ab}Means with dissimilar superscripts in a row are significantly different ($p < 0.05$)

Table 2. Effect of slaughter age on carcass characteristics.

Parameter	Slaughter age			P-value
	8 wk Mean \pm SD	10 wk Mean \pm SD	12 wk Mean \pm SD	
Live weight (g)	586.66 ^c \pm 39.45	741.94 ^b \pm 51.70	989.38 ^a \pm 64.85	$P < 0.001$
Dressing yield (%)	66.90 ^b \pm 1.50	64.53 ^c \pm 1.78	68.18 ^a \pm 1.43	$P < 0.001$

^{abc}Means with dissimilar superscripts in a row are significantly different ($p < 0.05$)

Table 3. Expected response to selection for 8 weeks body weight (g) in second generation (G₂).

Genotype	Sex	Population tested		Population selected		Selection Differential (S)	Selection Intensity (i)	Phenotypic standard deviation (sd)	Heritability of the trait (h ²)	Expected response to selection (R)
		No.	Aver.	No.	Aver.					
ND	M	420	497.32	85	599.94	102.63	4.73	21.68	0.499±0.030	51.21
	F	479	401.77	230	449.94	48.18	4.35	11.07		24.03
H	M	182	582.46	45	659.66	77.20	0.84	92.16	0.489±0.027	37.74
	F	186	519.78	115	551.43	31.65	1.64	19.30		15.47
NN	M	157	536.61	45	591.56	54.94	0.96	57.25	0.478±0.034	26.26
	F	168	428.07	115	448.27	20.19	0.43	46.62		9.65

Table 4. Comparison between second generation (G₂) and foundation stock (G₀).

Parameter	Generation	Genotype		
		ND	H	NN
Chick wt (g)	G ₀	27.74	28.00	24.96
	G ₂	25.77	26.06	25.40
	gain/change	-1.97	-1.94	0.44
8 wk wt (g)	G ₀	342.20	375.17	331.64
	G ₂	449.54	551.12	482.34
	gain/change	107.34	175.95	150.70
ASM (days)	G ₀	160.73	166.06	160.04
	G ₂	150.27	158.27	141.29
		-10.465	-7.79	-18.75
40 wk wt (g)	G ₀	1177.35	1377.53	1170.47
	G ₂	1380.26	1714.89	1243.29
	gain/change	202.91	337.36	72.82
Egg Prod. (no.)	G ₀ (24-40 wks)	60.04	56.19	55.28
	G ₂ (24-40 wks)	74.02	58.33	66.86
	gain/change	13.98	2.14	11.58
Egg wt (g)	G ₀ (at 38-40 wks)	42.26	41.14	41.61
	G ₂ (at 38-40 wks)	41.78	42.61	38.37
	gain/change	-0.48	1.47	-3.24

4. Conclusions

The results revealed that among the indigenous genotypes H genotype was superior in terms of body weight, and NN genotype was for dressing percentage. NN genotype was also found reaching maturity earlier but attaining a lighter mature weight. In each generation slight response was obtained for selection. These findings give an impetus for continuing the pure breeding research for more generations.

Conflict of interest

None to declare.

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