

Original Article

Influence of the type of breed and line of agricultural poultry on the morphological composition of eggs

Influência do tipo de raça e linhagem de aves agrícolas na composição morfológica dos ovos

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Abstract

The article presents, for the first time, the results of differentiating chickens, quails, turkeys, and ducks based on the morphological characteristics of their eggs and establishes the influence of species, breed, and line affiliation on egg morphological parameters. The process of agglomeration of breeds and lines of birds occurred within species affiliation, which in turn was influenced by differences in the morphological structure of eggs caused by the direction of selection. In quails and ducks, the grouping of breeds and lines occurred more rapidly than in chickens and turkeys, which is due to the intensity of selection for desirable traits. Species affiliation determined the differences in the relative weights of the shell, albumen, and yolk ($\eta^2 = 0.420-0.467$, $p = 0.000$), as well as the relative weights of the outer liquid and dense layers of the albumen ($\eta^2_{\text{ROLLA}} = 0.399$ and $\eta^2_{\text{RODLA}} = 0.351$, $p = 0.000$). Although quails and ducks were similar in the relative weight of albumen, they exhibited differences in its structural characteristics.

Keywords: chickens, quails, ducks, turkeys, morphological composition of the egg, species, breed, line, hierarchy analysis method.

Resumo

O artigo apresenta, pela primeira vez, os resultados da diferenciação de galinhas, codornas, peruas e patas com base nas características morfológicas de seus ovos, e estabelece a influência da espécie, raça e linhagem nos parâmetros morfológicos dos ovos. O processo de aglomeração de raças e linhagens de aves ocorreu dentro da mesma espécie, a qual, por sua vez, foi influenciada por diferenças na estrutura morfológica dos ovos causadas pela direção da seleção. Em codornas e patas, o agrupamento de raças e linhagens ocorreu mais rapidamente do que em galinhas e peruas, o que se deve à intensidade da seleção para características desejáveis. A afiliação de espécies determinou as diferenças nos pesos relativos da casca, do albúmen e da gema ($\eta^2 = 0,420 - 0,467$, $p = 0,000$), bem como nos pesos relativos do líquido externo e das camadas densas do albúmen ($\eta^2_{\text{ROLLA}} = 0,399$ e $\eta^2_{\text{RODLA}} = 0,351$, $p = 0,000$). Embora ovos de codornas e patas fossem semelhantes no peso relativo de albúmen, exibiram diferenças em suas características estruturais.

Palavras-chave: galinhas, codornas, patas, peruas, composição morfológica do ovo, espécie, raça, linhagem, método de análise de hierarquia.

1. Introduction

In a relatively short period, scientific and technological progress has transformed poultry farming into a fully developed branch of animal husbandry, now leading in the global production of animal protein for human nutrition. This is primarily due to the creation of numerous breeds of agricultural poultry, many of which have industrial significance (Saxena and Kolluri, 2018; Thiruvengadan and Prabakaran, 2017).

Eggs of domestic birds of different species show significant variation in morphological composition, owing to differing evolutionary processes; however, the range of variability in the relative weight of egg components is quite narrow. Depending on the bird species, the shell accounts for 9–12% of the total egg weight, the albumen approximately 60%, and the yolk 30–33% (Nys and Guyot, 2011; Birkhead et al., 2018). Y. Singh et al. found statistically

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significant differences in morphological qualities (shell, albumen, and yolk weight) between the eggs of domestic ducks and chickens (Singh, et al., 2022; Duman et al., 2016). Sun et al. reported substantial differences in egg quality traits among domestic chickens, ducks, geese, turkeys, quails, and pigeons. The relative yolk weight varied by species within the range of 19.3–37.9% (Sun et al., 2019).

Within species, specialization of birds for priority traits has led to the emergence of pools with similar exterior and productivity characteristics, forming the basis for breed development and, through more targeted specialization, for lines. To varying degrees, poultry breeds within a single species differ in quantitative and qualitative traits, including the morphological structure of eggs. For instance, when comparing the maternal lines Hy-Line Brown (brown-shelled eggs) and Hy-Line W-36 White Leghorn (white-shelled eggs), significant differences in the relative shell mass were observed throughout the study period (32–85 weeks) ($p < 0.0001$). Differences in relative albumen and yolk mass were found only during certain age periods. Eggs from Hy-Line Brown hens differed from Messa 43 eggs by having a lower albumen proportion and a higher shell proportion (Sharma et al., 2022; Biesiada-Drzazga et al., 2020).

A multi-breed experiment with 25 commercial and traditional laying hen lines demonstrated a high degree of variability (> 0.8) in the mass of albumen, yolk, and shell, attributed to breed or line affiliation. The variability of these traits within lines of a single breed was much smaller than between breeds (Hocking et al., 2003). Importantly, selection intensity negatively impacts the sensory qualities of egg components, particularly the yolk (Zhang et al., 2022).

In ducks, flock P11 eggs contained 1.6% more albumen but 2.0% less yolk compared to flock P22 eggs. Statistically significant differences in density and pH were observed between the liquid and dense layers of albumen. Eggs from Cherry Valley ducks had a higher relative albumen mass, while those from Peking ducks had higher yolk and shell proportions. The albumen/yolk ratio was greater in Peking ducks compared to Cherry Valley ducks ($p < 0.01$) (Kokoszyski et al., 2007; Galić et al., 2019).

From a biological standpoint, the primary function of a bird's egg is to ensure normal embryogenesis and the emergence of viable offspring. The quantitative and qualitative components of eggs are interrelated, and changes in one component affect the others (Javůrková and Mikšík, 2023; Boleli et al., 2016).

Egg weight is strongly phenotypically correlated with egg shape, absolute and relative weights of shell, albumen, and yolk. Therefore, variations in egg weight influence the variability of its components (Wolc et al., 2012; Hristakieva, et al., 2017).

In North Caucasian Bronze turkeys, phenotypic correlations between egg weight and shell weight were observed at 34 and 46 weeks of age ($r = 0.51$ and $r = 0.50$; $p < 0.05$). In brown-shelled chicken eggs, albumen and yolk weights showed positive phenotypic correlations with egg and shell weights: albumen with egg and shell (0.94 and 0.32), yolk with egg and shell (0.71 and 0.31). In quails, egg weight was positively correlated with albumen

weight ($r = 0.88$; $p < 0.001$) and yolk weight ($r = 0.26$; $p < 0.001$). Variability in egg weight was more closely linked to variability in albumen weight (Zhang et al., 2005; Sari et al., 2016).

This conclusion is supported by research on the effects of protein-deficient diets on egg morphology. Protein-deficiency-related decreases in egg weight had less effect on yolk and shell weights than on total egg weight and albumen weight (Fisher, 1969).

Sinha B. et al., based on a synthesis of data, noted that shell, albumen, and yolk weights are breed- and line-specific in chickens, but the absolute and relative albumen mass most strongly reflects breed and line identity. The possibility of developing chicken lines characterized by dense albumen mass was suggested (Sinha et al., 2018).

In quails, phenotypic correlations between egg weight and egg component weights varied depending on breed. In dark brown quails, phenotypic correlation coefficients for egg weight with shell, yolk, and albumen mass were $r_{\text{shell}} = 0.17$ ($p > 0.5$), $r_{\text{yolk}} = 0.20$ ($p < 0.01$), $r_{\text{albumen}} = 0.62$ ($p < 0.01$). In light brown quails: $r_{\text{shell}} = 0.35$ ($p < 0.01$), $r_{\text{yolk}} = 0.61$ ($p < 0.01$), $r_{\text{albumen}} = 0.74$ ($p < 0.01$). In white quails: $r_{\text{shell}} = 0.67$ ($p < 0.01$), $r_{\text{yolk}} = 0.62$ ($p < 0.01$), $r_{\text{albumen}} = 0.68$ ($p < 0.01$). Regardless of breed, egg weight was most strongly associated with albumen mass (Hassan et al., 2017).

Thus, the structure of the egg reflects, to varying degrees, the species, breed, and line affiliation of the bird. The latter two are driven by artificial selection for specific desirable traits. Due to the rapid maturity, high reproductive rate, and adaptability of poultry, the shaping force of artificial selection is more pronounced in poultry breeding than in other livestock sectors.

The aim of our study was to investigate the similarities and differences between species, breeds, and lines of agricultural poultry under the influence of artificial selection for economically valuable traits.

2. Material and Methods

The study was conducted at the LLP “North Kazakhstan Agricultural Research Institute” (Republic of Kazakhstan) and the Siberian Research Institute of Poultry Farming — a branch of the FSBSI “Omsk Agricultural Scientific Center.”

The research object comprised eggs from flocks of quails (*Coturnix coturnix*) of the Japanese, Omsk, Pharaoh, and Texas White breeds; chickens (*Gallus gallus*) from the maternal lines of Dekalb White, Hisex Brown, and Ross 308 crossbreeds; ducks (*Anas platyrhynchos*) of the Agidel, Ansar, and Star lines; and turkeys (*Meleagris gallopavo*) of the maternal line of the Hybrid Converter crossbreed.

For each breed and line, 120 eggs laid on the same day were studied. The birds were aged 52 weeks for chickens, ducks, and turkeys, and 40 weeks for quails.

The procedure sequentially involved determining the weight of the whole egg (ME), followed by measuring the absolute weights of the shell (MS), albumen (MA), and yolk (MY). The albumen was further separated into the following layers with their respective weights recorded: outer liquid albumen (MOLLA), outer dense albumen (MODLA), inner

liquid albumen (MLDLA), and inner dense albumen (MILLA). All weights were measured with an accuracy of 0.01 g. Shell thickness was measured to an accuracy of 0.01 mm.

Based on the obtained data, the following relative masses were calculated:

Relative shell mass ($RS = MS/ME \times 100, \%$),

Relative albumen mass ($RA = MA/ME \times 100, \%$),

Relative yolk mass ($RY = MY/ME \times 100, \%$),

Relative outer liquid albumen ($ROLLA = MOLLA/MA \times 100, \%$),

Relative outer dense albumen ($RODLA = MODLA/MA \times 100, \%$),

Relative inner liquid albumen ($RILLA = MLDLA/MA \times 100, \%$),

Relative inner dense albumen ($RIDLA = MILLA/MA \times 100, \%$).

Statistical analysis was performed using SPSS software, version 23. For the measured parameters, means (M) and their standard errors ($\pm$ SEM) were presented. The degree of variability in traits was assessed using the coefficient of variation (Cv). Species, breed, and line affiliation of the birds were considered as factors influencing the morphological composition of the eggs. The strength of factor influence (η^2) was evaluated using analysis of variance (ANOVA). Statistical significance was assessed using Fisher's F-test, with significance levels set at $p < 0.05$ – 0.001 .

The degree of correlation between varying traits was determined by phenotypic correlation coefficients. The intraclass correlation coefficient (ICC) was used to quantify the degree of trait similarity among breeds and lines.

Data on the relative weights of shell, yolk, albumen, and albumen fractions were used for hierarchical classification of the studied species, breeds, and lines. Based on these data, a dendrogram was constructed, and cluster similarity was assessed by the squared Euclidean distance.

3. Results and Discussion

The egg mass corresponded to the recommended values for birds of the respective age. The coefficients of

variation for egg mass across all breeds and lines were low ($Cv < 10\%$), indicating consolidation of the flock with respect to this trait (Table 1).

Statistically significant interspecies differences were observed in the relative mass of albumen and yolk. The most pronounced differences in these parameters were identified between ducks and the other three avian species: ducks differed from chickens, quail, and turkeys (all belonging to the order Galliformes) by having a lower albumen mass and a higher yolk mass. Specifically, the relative albumen mass in duck eggs was lower by 6.59% compared to quail, 7.41% compared to chickens, and 8.53% compared to turkeys, while the relative yolk mass was correspondingly higher by 5.05%, 7.37%, and 8.42%, respectively. Interestingly, as the egg mass increases among Galliformes species (from quail $\rightarrow$ chickens $\rightarrow$ turkeys), the disparity in relative albumen and yolk mass compared to duck eggs also increases. Quail, characterized by smaller egg size, exhibited a lower relative shell mass compared to other bird species by 1.43–1.54%. In contrast, the differences in relative shell mass between ducks, chickens, and turkeys were minimal (Table 2).

Statistically significant differences between quail and chickens, ducks, and turkeys were observed in the relative mass of the albumen layers. Compared to other bird species, quail eggs had a higher proportion of outer thin albumen and a lower proportion of outer thick and inner albumen layers. Chickens and turkeys also differed significantly from ducks in the relative proportions of albumen layers. Specifically, chickens exhibited a higher proportion of both thick albumen layers and a lower proportion of both thin albumen layers compared to ducks. Statistically significant differences between chickens and turkeys were found in the relative mass of the outer thin, inner thin, and inner thick albumen layers. The relative mass of the outer thick albumen layer was nearly the same between these two species. Between ducks and turkeys, significant differences were noted in the relative mass of the outer albumen layers and the inner thin albumen layer. Ducks had a greater proportion of outer thin albumen, whereas chickens had more outer thick albumen.

Table 1. Egg Mass and Coefficient of Variation ($n = 120$).

Type	Breed / Line	M $\pm$ SEM, g	Cv, %
Quail	Japanese	12.78 $\pm$ 0.111	7.66
	Omsk	13.56 $\pm$ 0.102	8.54
	Pharaoh	13.74 $\pm$ 0.100	7.68
	Texas White	15.31 $\pm$ 0.068	8.61
Chicken	Dekalb White	64.12 $\pm$ 0.357	8.85
	Hisex Brown	65.22 $\pm$ 0.334	9.40
	Ross 308	67.78 $\pm$ 0.811	6.53
Domestic Duck	Agidel Line	74.93 $\pm$ 0.752	7.54
	Ansar Line	75.41 $\pm$ 0.640	7.86
	Star Line	77.25 $\pm$ 0.693	6.56
Turkey	Hybrid Converter	95.83 $\pm$ 0.821	7.33

Table 2. Relative Mass of Egg Components (n = 120), %.

Type	Breed / Line Shell		Mass of Egg Components			Mass of Outer Albumen Layers		Mass of Inner Albumen Layers	
			Albumen	Yolk	liquid	dense	liquid	dense	RIDLA
			RS	RA	RY	ROLLA	RODLA	RILLA	
Quail	Japanese	M	10.31 ^{a2a3a4}	57.98 ^{a2a3a4}	31.71 ^{a3a4}	41.21 ^{a2a3a4}	36.66 ^{a3a4}	20.35 ^{a2}	1.77
		SEM	0.073	0.275	0.323	1.195	0.664	0.791	0.042
	Omsk	M	9.42 ^{a1a4}	59.07 ^{a1a3a4}	31.52 ^{a3a4}	44.32 ^{a1a3a4}	36.22 ^{a3a4}	17.72 ^{a1a3a4}	1.73 ^{a4}
		SEM	0.065	0.262	0.244	0.644	0.495	0.373	0.092
	Pharaoh	M	9.54 ^{a1}	60.14 ^{a1a2}	30.13 ^{a1a2}	36.54 ^{a1a2}	41.58 ^{a1a2}	19.92 ^{a2}	1.96
		SEM	0.104	0.378	0.364	0.897	0.900	0.803	0.173
	Texas White	M	9.63 ^{a1a2}	59.98 ^{a1a2}	30.39 ^{a1a2}	36.46 ^{a1a2}	41.77 ^{a1a2}	19.79 ^{a2}	1.97 ^{a2}
		SEM	0.059	0.151	0.147	0.555	0.537	0.337	0.071
	Species Average	M	9.69 ^{bcd}	59.29 ^{bcd}	31.02 ^{bcd}	40.10 ^{bcd}	38.83 ^{bcd}	19.22 ^{bcd}	1.85 ^{bcd}
		SEM	0.039	0.131	0.128	0.431	0.330	0.259	0.045
Chicken	Dekalb White	M	10.99 ^{b2}	63.11 ^{b3}	25.90 ^{b2b3}	30.54 ^{b2b3}	44.02 ^{b2b3}	23.40 ^{b2b3}	2.04 ^{b2b3}
		SEM	0.075	0.166	0.165	0.732	0.533	0.486	0.069
	Hisex Brown	M	11.87 ^{b1b3}	63.54 ^{b3}	24.59 ^{b1b3}	28.11 ^{b1b3}	48.34 ^{b1b3}	20.27 ^{b1}	3.28 ^{b1b3}
		SEM	0.107	0.260	0.215	0.501	0.480	0.251	0.056
	Ross 308	M	10.81 ^{b2}	56.12 ^{b1b2}	33.07 ^{b1b2}	23.78 ^{b1b2}	53.23 ^{b1b2}	20.10 ^{b1}	2.89 ^{b1b2}
		SEM	0.075	0.196	0.171	0.420	0.495	0.300	0.035
	Species Average	M	11.23 ^a	60.87 ^{acd}	27.90 ^{acd}	27.41 ^{acd}	48.63 ^{ac}	21.21 ^{acd}	2.75 ^{acd}
		SEM	0.056	0.220	0.227	0.354	0.351	0.219	0.042
Domestic Duck	Agidel Line	M	10.93 ^{c2c3}	53.83	35.24	29.22 ^{c2c3}	41.30 ^{c2c3}	26.96 ^{c2c3}	2.52 ^{c2c3}
		SEM	0.047	0.381	0.429	0.407	0.353	0.241	0.022
	Ansar Line	M	11.51 ^{c1}	53.36	35.13	36.06 ^{c1}	43.30 ^{c1}	19.10 ^{c1}	1.54 ^{c1}
		SEM	0.056	0.212	0.232	0.262	0.241	0.290	0.044
	Star Line	M	11.54 ^{c1}	52.95	35.51	35.27 ^{c1}	43.53 ^{c1}	19.62 ^{c1}	1.58 ^{c1}
		SEM	0.067	0.321	0.335	0.236	0.329	0.369	0.058
	Species Average	M	11.27 ^a	53.46 ^{abd}	35.27 ^{abd}	32.96 ^{abd}	42.51 ^{abd}	22.57 ^{abd}	1.96 ^{ab}
		SEM	0.035	0.19	0.214	0.262	0.195	0.256	0.033
Turkey	Hybrid Converter	M	11.16 ^a	61.99 ^{abc}	26.85 ^{abc}	24.29 ^{abc}	49.52 ^{ac}	24.05 ^{abc}	2.14 ^{ab}
		SEM	0.076	0.240	0.247	0.281	0.499	0.278	0.074

Note: 1) Statistically significant differences between species: a – compared to quail, b – compared to chicken, c – compared to duck, d – compared to turkey; 2) Statistically significant differences between breeds/lines: Quail: a1 – compared to Japanese, a2 – compared to Omsk, a3 – compared to Pharaoh, a4 – compared to Texas White; Chicken: b1 – compared to Dekalb White line, b2 – compared to Hysex Brown line, b3 – compared to Ross 308 line; Ducks: c1 – compared to Agidel line, c2 – compared to Ansar line, c3 – compared to Star line.

Thus, statistically significant differences were established among all four bird species in terms of the relative mass of yolk, albumen, and the inner and outer thin albumen layers. In quail, chickens, and turkeys—all belonging to the order Galliformes—statistically significant correlations were identified between the relative mass of total albumen and the relative mass of its outer layers: a positive correlation with the outer thin albumen ($r_{\text{quail}} = 0.155$, $r_{\text{chicken}} = 0.355$, $r_{\text{turkey}} = 0.121$; $p < 0.05$ – 0.01) and a negative correlation with the outer thick albumen ($r_{\text{quail}} = -0.116$, $r_{\text{chicken}} = -0.422$, $r_{\text{turkey}} = -0.148$; $p < 0.05$ – 0.01). The

correlation coefficients for both albumen fractions were higher in chickens than in the other two species. Conversely, in ducks, a negative correlation was found between the relative mass of total albumen and the outer thin layer ($r_{\text{duck}} = -0.412$, $p < 0.01$), and a positive correlation with the outer thick layer ($r_{\text{duck}} = 0.409$, $p < 0.01$). Across all bird species, common trends in albumen fraction correlations were observed: a negative correlation between the relative mass of the outer thin and thick albumen layers ($r_{\text{quail}} = -0.789$, $r_{\text{chicken}} = -0.816$, $r_{\text{turkey}} = -0.853$, $r_{\text{duck}} = -0.299$; $p < 0.01$), and between the

outer thick and inner thin albumen layers ($r_{\text{quail}} = -0.222$, $r_{\text{chicken}} = -0.337$, $r_{\text{turkey}} = -0.867$, $r_{\text{duck}} = -0.426$; $p < 0.05$ – 0.01). A negative correlation between the inner thin and thick layers was identified in quail, chickens, and turkeys ($r_{\text{quail}} = -0.145$, $r_{\text{chicken}} = -0.221$, $r_{\text{turkey}} = -0.141$; $p < 0.05$ – 0.01), while in ducks, this correlation was positive ($r_{\text{duck}} = 0.549$; $p < 0.05$ – 0.01).

Differences between breeds were attributed to selection goals for specific productivity traits. Among quail, the Japanese breed (egg-type) differed from the Omsk breed (dual-purpose) and the meat-type Pharaoh and Texas White breeds in the relative mass of shell, albumen, and the outer thin albumen layer. A trend was observed: quail of the Japanese breed, which had smaller egg mass compared to other breeds, exhibited a higher relative shell mass. This was confirmed by a moderate intraclass correlation coefficient ($\text{ICC} = 0.315$, $p = 0.035$). Omsk quail differed from the other three breeds in total albumen mass and in the mass of both thin albumen fractions—outer and inner. The Pharaoh and Texas White breeds were similar in the relative mass of shell and yolk, and differed from the Japanese and Omsk breeds in the relative mass of albumen and its outer layers (outer thin and outer thick). Additionally, these meat-type breeds differed from the Japanese and Omsk breeds in the relative mass of the inner thin albumen layer.

Among chickens, statistically significant differences were found between all three crosses in the relative mass of yolk, outer thin and thick albumen layers, and the inner thick albumen layer. Eggs from the white-shelled cross Dekalb White differed from those of the brown-shelled crosses Hisex Brown and Ross 308 by having a higher relative mass of the inner thin albumen layer. This feature appeared to be a consistent pattern, as evidenced by a statistically significant but weak correlation between cross type and the relative mass of the inner thin albumen layer ($\text{ICC} = 0.198$, $p = 0.045$).

Hisex Brown eggs had a higher relative shell mass. The relative shell mass of the chicken crosses increased in the following order: Ross 308 → Dekalb White → Hisex Brown. The same order was observed for shell thickness: Ross 308 (0.35 ± 0.002 mm) → Dekalb White (0.37 ± 0.002 mm) → Hisex Brown (0.38 ± 0.003 mm). This pattern suggests a potential interdependence between these parameters. A positive correlation was identified between relative shell mass and shell thickness, with a strong correlation in the Hisex Brown cross ($r = 0.800$, $p < 0.01$), moderate in Dekalb White ($r = 0.413$, $p < 0.01$), and Ross 308 ($r = 0.376$, $p < 0.01$). Notably, the strength of these correlations followed the same order as the increasing trends in shell mass and thickness.

The albumen-to-yolk ratio in eggs from the Ross 308 broiler cross was 1.74, compared to 2.62 and 2.49 in the Hisex Brown and Dekalb White layer crosses, respectively. In egg-type chickens, selection is primarily focused on egg-laying performance, while in meat-type chickens, emphasis is placed on growth rate and hatchability. It was found that eggs from the Ross 308 broiler cross differed from those of the egg-laying crosses (Hisex Brown and Dekalb White) by having a lower relative albumen mass and a higher relative yolk mass. Statistically significant

intraclass correlation coefficients indicated an association between these traits and the productivity direction ($\text{ICCRA} = 0.114$, $p = 0.049$; $\text{ICCRY} = 0.219$, $p = 0.039$). This association was slightly stronger for relative yolk mass than for albumen mass.

The Ansar and Star duck lines were comparable in productivity: their average live weights were 3.5 kg and 3.7 kg, and egg production over 40 weeks totaled 253 and 258 eggs, respectively. These lines were also similar in egg structural composition. In contrast, the Agidel line, characterized by lower live weight (3.2 kg) and egg production (220 eggs), differed from Ansar and Star in the relative mass of the shell and albumen layers.

Shell thickness in Agidel duck eggs was 0.30 ± 0.001 mm, compared to 0.35 ± 0.002 mm and 0.34 ± 0.002 mm in the Ansar and Star lines, respectively, closely mirroring differences in shell mass. However, when analyzed separately by line, a statistically significant correlation between shell thickness and relative shell mass was found only in the Ansar line ($r = 0.392$, $p < 0.01$). Although the lines were comparable in total relative albumen mass, the Agidel line differed from the Star and Ansar lines in the relative mass of individual albumen layers. The most pronounced differences were observed in the liquid fractions. Eggs from the Agidel line had a lower relative mass of the outer thin albumen layer by 6.84% and 6.05%, respectively, compared to the Ansar and Star lines. However, this was compensated by a higher relative mass of the inner thin albumen layer in Agidel eggs—by 7.86% and 7.34%, respectively. The relative mass of the outer thick albumen layer was also lower in the Agidel line by 2.00% and 2.33%.

Comparing the structurally and productively similar duck lines (Ansar and Star) to the distinct Agidel line enabled the identification of a significant relationship between egg structure and productivity traits: $\text{ICCRS} = 0.114$, $p = 0.049$; $\text{ICCRolla} = 0.152$, $p = 0.019$; $\text{ICCRDLA} = 0.111$, $p = 0.034$; $\text{ICCRILLA} = 0.122$, $p = 0.024$; $\text{ICCRIDLA} = 0.222$, $p = 0.009$.

The effect of the “species” factor on all studied traits was statistically significant. Its influence on the relative mass of shell, albumen, and yolk was of moderate strength and similar in magnitude across these traits. In terms of albumen layer distribution, species had a moderate effect on the outer layers and a weaker effect on the inner layers. The influence of the “breed/line” factor was statistically significant in most cases. Notable species-specific differences were observed in how breed influenced the relative mass of egg components. In quail, the breed effect was most prominent for shell mass and the mass of the outer albumen layers (weak influence). A moderate effect of the “breed” factor was found for outer thin and inner albumen layers, and a weak effect for shell mass. In chickens, the magnitude of the “line” effect was more pronounced than in quail or ducks, indicating a higher degree of specialization in the selected lines and maternal forms regarding productivity traits. Chicken lines differed more substantially in albumen and yolk mass (with significant and strong effects, respectively), and also in the mass of the outer thin and inner thick albumen layers. For other parameters, the influence was weak but remained statistically significant (Table 3).

The greater influence of bird species and breed/line on the variability of the relative mass of the outer thin albumen layer compared to the outer thick layer is consistent with the findings of Y. Wan, who reported heritability coefficients of 0.39 and 0.31 for these respective albumen layers (Wan et al., 2019).

At the initial stage of agglomerative clustering, the meat-type quail breeds Pharaoh and Texas White (Saxena and Kolluri, 2018) were grouped into a single cluster, reflecting their similarity across all parameters analyzed in the study. In the second stage, the duck lines Ansar and Star (Hocking et al., 2003) were clustered together, as they also exhibited close similarities. The third stage involved clustering of the Omsk and Japanese quail breeds (Nys and Guyot, 2011), which showed similar values for

the relative masses of yolk and both outer and inner thick albumen layers. At the fourth stage, the Dekalb White and Hisex Brown chicken lines (Sun et al., 2019) were grouped together due to their similarity in relative albumen mass.

Interestingly, in the fifth stage of agglomeration, the cluster formed by the egg-type lines Dekalb White and Hisex Brown merged not with the Ross 308 chicken line, but with the turkey group. This was explained by the fact that the egg-type chicken cluster exhibited less dissimilarity from the turkey cluster in terms of relative albumen and yolk mass ($\eta^2_{RA} = 0.053$, $\eta^2_{RY} = 0.086$, $p = 0.000$) and the mass of the outer thin and thick albumen layers ($\eta^2_{ROLLA} = 0.057$, $\eta^2_{RODLA} = 0.094$, $p = 0.000$), compared to the meat-type Ross 308 chicken line cluster ($\eta^2_{RA} = 0.689$, $\eta^2_{RY} = 0.763$; $\eta^2_{ROLLA} = 0.155$, $\eta^2_{RODLA} = 0.253$, $p = 0.000$) (Figure 1).

Table 3. Influence of Species and Breed on the Variability of the Relative Mass of Egg Components (n = 120).

Factor Shell		Egg Component Mass			Outer Albumen Layer Mass		Inner Albumen Layer Mass	
		Albumen	Yolk	liquid	dense	liquid	dense	RIDLA
Type		RS	RA	RY	ROLLA	RODLA	RILLA	
Breed/Line	η^2	0.436	0.467	0.420	0.399	0.351	0.096	0.202
	p	0.000	0.000	0.000	0.000	0.000	0.000	0.000
Quail	η^2	0.184	0.096	0.066	0.175	0.173	0.047	0.017
	p	0.000	0.000	0.000	0.000	0.000	0.000	0.101
Chicken	η^2	0.200	0.691	0.779	0.324	0.177	0.133	0.431
	p	0.000	0.000	0.000	0.000	0.000	0.000	0.000
Duck	η^2	0.183	0.009	0.001	0.406	0.075	0.581	0.550
	p	0.000	0.190	0.799	0.000	0.000	0.000	0.000

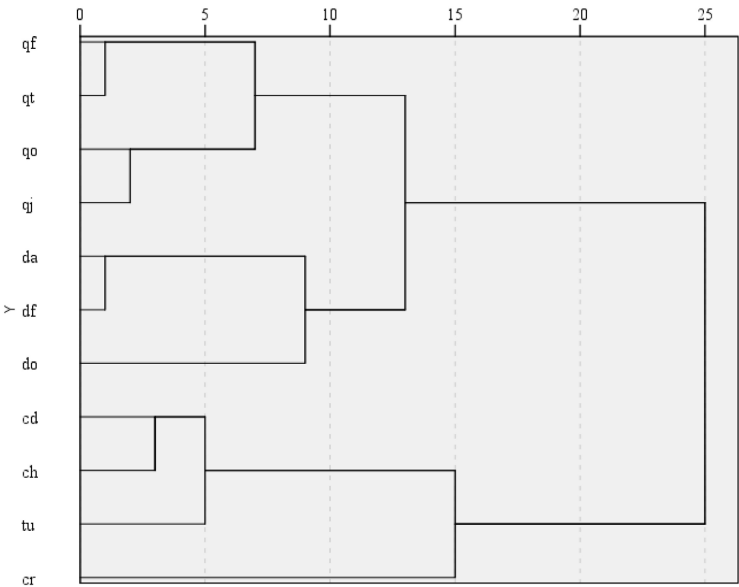


Figure 1. Hierarchical Analysis of Morphological Traits of Poultry Species and Breeds. qf – Pharaoh quail breed, qt – Texas White quail breed, qo – Omsk quail breed, qj – Japanese quail breed; da – Ansar duck line, df – Star duck line, do – Agidel duck line; cd – Dekalb White chicken line, ch – Hisex Brown chicken line, cr – Ross 308 chicken line; tu – Hybrid Converter turkey line.

The aggregation of breeds into species-level clusters occurred earlier for quail and ducks than for chickens. Moreover, quail and ducks—despite belonging to different avian orders—merged into a single cluster prior to the clustering of the Ross 308 meat-type chicken line with egg-type chickens and turkeys. This pattern is attributed to the more pronounced differences in relative masses of the outer thin and outer thick albumen layers between chickens and quail ($\eta^2_{\text{ROLLA}} = 0.304$ and $\eta^2_{\text{RODLA}} = 0.278$, $p = 0.000$), as well as between chickens and ducks ($\eta^2_{\text{ROLLA}} = 0.214$ and $\eta^2_{\text{RODLA}} = 0.263$, $p = 0.000$). Although statistically significant, the differences between quail and ducks were of minimal magnitude ($\eta^2_{\text{ROLLA}} = 0.069$ and $\eta^2_{\text{RODLA}} = 0.076$, $p = 0.000$).

These findings are consistent with previous studies, which reported that selection with a narrow focus on productivity traits has led to greater genetic and phenotypic differentiation among chicken lines compared to other poultry species. Furthermore, industrial breeds and lines tend to differ more significantly from local or traditional breeds (Vaarst et al., 2015).

The scientific classification of poultry species, breeds, and lines provides a systematic framework for organizing objects, generating new generalizations and hypotheses, and analyzing relationships both within and between classes.

The results of our study demonstrated that the agglomeration process among poultry breeds and lines followed species boundaries, which were, in turn, influenced by morphological differences in egg structure associated with selective breeding objectives. Breed aggregation within quail and duck species occurred more rapidly than in chickens and turkeys, reflecting differences in the intensity and focus of selection for desirable traits.

Species affiliation significantly determined differences in the relative mass of eggshell, albumen, and yolk ($\eta^2 = 0.420\text{--}0.467$, $p = 0.000$), as well as in the relative mass of the outer thin and thick albumen layers ($\eta^2_{\text{ROLLA}} = 0.399$ and $\eta^2_{\text{RODLA}} = 0.3516$, $p = 0.000$). While quail and ducks were similar in terms of total relative albumen mass, they differed in the structural composition of the albumen.

4. Conclusions

This study demonstrated that the morphological structure of eggs is significantly influenced by species, breed, and line affiliation. Species had a moderate effect on the relative mass of the shell, albumen, and yolk ($\eta^2 = 0.420\text{--}0.467$, $p = 0.000$), and on the outer albumen layers ($\eta^2_{\text{ROLLA}} = 0.399$, $\eta^2_{\text{RODLA}} = 0.351$, $p = 0.000$). While quail and ducks showed similar total albumen mass, they differed in albumen layer distribution. Breed and line effects were especially notable in chickens. Egg-type lines (Dekalb White, Hisex Brown) had higher albumen-to-yolk ratios (2.49–2.62) than the meat-type Ross 308 (1.74), with yolk mass more strongly associated with productivity direction (ICCRY = 0.219, $p = 0.039$). In ducks, the Agidel line showed structural shifts in albumen distribution compared to the Ansar and Star lines, despite similar total albumen mass. Hierarchical clustering revealed earlier aggregation of

quail and duck breeds compared to chickens, indicating stronger breed divergence in chickens due to selection intensity. Egg morphology thus reflects both species identity and breeding objectives, making it a valuable tool for classification and breeding program assessment.

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Data Availability Statement

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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