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Multiple Sequence Alignment and Phylogenetic Analysis of *Growth Differentiation Factor 9* Gene for Egg Production in Poultry Species

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ABSTRACT

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Bioinformatic analysis Evolutionary tree DNA sequences Growth differentiation factor 9 gene (*GDF9*) is an oocyte-secreted factor which controls the functions of the ovarian cells, regulates the development and caliber of the egg. *GDF9* DNA and protein sequences from NCBI were used to computationally investigate molecular genetic variation in poultry species. Results showed that percentage similarities of DNA and protein sequences ranged from 78.45 to 92.80% and 83.63 to 94.74%, respectively. The phylogenetic analysis results showed that chicken and turkey were closely related. The high percent similarities and close relatedness of poultry species show that *GDF9* can be used as a candidate gene for improving egg production in poultry species.

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Introduction

Poultry species refers to a group of birds which include chicken, turkey, Ostrich and they are bred for their ability to produce products including eggs (FAO, 2022). According to Liu et al. (2019), egg production plays an essential role in food security and alleviation of unemployment. However, due to the ongoing poor performance in egg production, these needs are barely met (Beesabathuni et al., 2018). This means that there is a constant need to improve egg production performance. Growth differentiation factor 9 gene (GDF9) is an oocyte-secreted factor that is responsible for controlling the functions of the ovarian cells as well as regulating the development and quality of the egg, subsequently making it a good candidate gene for improving egg production performance (Belli and Shimasaki, 2018). According to Zamani et al. (2015), studying genes through bioinformatics allows for determination of genetic similarities and variation in their sequences in a faster manner thus allowing researchers to know how these findings affect the production performance of species. Multiple sequence alignment is essential in bioinformatics for identifying homologous regions on biological sequences, such as DNA, and it is a crucial prerequisite for phylogenetic tree construction (Bodenhofer et al., 2015). Phylogenetic analysis is a simple bioinformatic method that aid in mapping the lineage of genes across various species, as well as predicting the interactions between the species based on their evolutionary relationship (Singh et al., 2023). According to Pramod et al. (2019), phylogenetic analysis can provide important data that may assist scientist with implementing effective animal management and conservation strategies. In a scientific study, Egom et al. (2019), was able to document that the percentage identity and similarities of insulin-like growth factor-1 gene in poultry species fell in a range of 86-99% and was able to conclude that the gene was highly effective for its protein function improvement.

Furthermore, Karthikeyan *et al.* (2020), documented that the Osteopontin gene in Bos taurus exhibited 99% similarity with the closely related ruminant species. This proves the ability to use bioinformatics for gene comparison within and among species. Additionally, Ewuola *et al.* (2018), has proved that bioinformatics methods, such as multiple alignment and phylogenetic analysis, are viable strategies to analyze variations and similarities in genes of poultry species. However, similarities and differences of *growth differentiation factor 9* gene in poultry species are not yet known. Hence the objective of the current study was to computationally investigate the molecular genetic variation of *growth differentiation factor 9* gene in poultry species. Findings from the current study will provide genetic variation information on percentage similarities of *growth differentiation factor 9* gene in poultry species which may be beneficial to scientists and researchers interested in improving egg production performance of these poultry species.

Materials and methods

Retrieval of deoxyribonucleic acid (DNA) and protein sequences

DNA and protein sequences of the *growth differentiation factor 9* gene of chicken, turkey and ostrich and their accession numbers were retrieved from the National Centre for Biotechnology Information (NCBI), as shown in Table 1.

Table 1. Retrieved DNA sequences of *growth differentiation factor 9* gene of selected poultry species with their accession numbers and sequence lengths

Species	DNA sequence accession number	Base pair	Protein seq. Accession number	Amino acid sequence length
Chicken	NM_206988.3	1707	NP_996871.3	457
Turkey	XM_003210484.4	1903	XP_003210532.1	457
Ostrich	XM_009678924.1	1341	XP_009677219.1	446

Alignment of multiple deoxyribonucleic acid (DNA) and protein sequences

Alignment of the multiple DNA and protein sequences were done using CLUSTAL-W software incorporated by Online Analysis Tools. Through this software utilization, a pairwise comparison was conducted and percentage similarities of all the retrieved sequences were obtained.

Phylogenetic analysis

Evolutionary relationships using the sequences of *growth differentiation factor* 9 gene retrieved from NCBI was conducted using Molecular Evolution and Genetic Analysis (MEGA) software, version 11. The phylogenetic tree was constructed using this software. A scale length of 0.05 was used for the construction of the tree.

Results

Retrieved of deoxyribonucleic acid (DNA) and protein sequences

Table 1 shows the accession, base pair and amino acid length numbers of DNA and protein sequences of the *growth differentiation factor 9* gene of chicken, turkey and ostrich which were retrieved from the National Centre for Biotechnology Information (NCBI). The *growth differentiation factor 9* gene and protein sequences of chicken, turkey and ostrich ranged from 1341 to 1903 bp and 446 to 457 bp, respectively.

Multiple alignment of DNA sequences

Figure 1 shows *growth differentiation factor 9* gene sequences of chicken, turkey and ostrich which were aligned using CLUSTAL-W incorporated by Online Analysis Tools. The percentage similarity of chicken and turkey DNA sequences was found to be 78.45% and 92.80% between chicken and ostrich. On the other hand, turkey and ostrich shared 83% similarity in their *GDF9* gene sequences.

Chicken Turkey Ostrich	ACGTGCGAAGACTCAGCCCGGCCGCACGCACACGTGGGTTCCACAGCCAGC
OSCIPICII	
Chicken Turkey	CTTTGCCTTTGGTTTCTAATCACAAAGAAATCGCACACAGCTGAAACGTATATAAGGGCT
Ostrich	
Chicken Turkey Ostrich	GAATGCTTAGGGGAGGCGTTGTTAGGCTACAGCC CGTGCTCGCATGCTCTTTTTCTCTCGAATGTTTTGGGGAGGCGTTGTTAGGCTGCAGCT
Chicken Turkey Ostrich	CTGAAGAGCATGGGGGGTACGTGGAGGATCTGTGTTTGTT
Chicken Turkey Ostrich	CTTTCTTCCAGCATCCAGTGCTCCCCCCGCTCCAGGGATCGCACAGCACCTGACAAGGTC CTTTCTTCTAGCATCCAGTGCTCCCCCCGCTCCAGGGATCGCGCAGCACCTGACAAACTC CTTTCTTCTAGCATCCAGTGTTCCCCCCGCTCCAGGGATCATGTAGTGTCTGACAAGACC ******* ****************************
Chicken Turkey Ostrich	TCCGGGCTCCTGGGAGCTCCTGAGCTCAGCGAGCTCCACCCGCTGCTGCGGCTG TCCGGGCTCCTGGAAGCTCCTGAGCTCAGCGAGCTCCACCCGCTGCTGCGGCTG TCCGGGTTACTGGTAGCTCCTGAGGACTACACCAGTAAGCTAAATCCGTTATTGCGGCTT ****** * **** **********************
Chicken Turkey Ostrich	CCCAAGGGCGTGAGCCGAGGCTATGCGCTCCTGCCCCCCTGCTCGAAGTGCTGTCTGAC CCCCAGGGTGCGAGCCGAGGCTGTGCCCTCCTGCCCCCCTGCTCAAAGTGCTGTCCGAC CCAAAAGGTGTGAGACGTGGATATGCCCTCCTGCCTCCCCTTCTCAAGGTGCTGTCTGAC ** * ** * *** * ** * ** * *** ****** *** *
chicken Turkey Ostrich	CACAGGACCCGATTACAGGAGACTTTCACTCGGTGGATTTGCTCTTCAACCTGGATCGT CACAGGGATCTGATTACAGGAGACCTTCACTCGGTGGATTTGCTCTTCAACCTGGATCGT CACAGGGACCTAATGAAAGGAGACCTTCACTCGGTGGATTTGCTCTTCAACCTGGATCGT ******* * * * ******* ***************
Chicken Turkey Ostrich	GTTACTGCTCTCGAGCACTTACTCAAGTCTGTCTTGCTCTACTCGTTTGACACCTCAGTT GTTACTGCTCTCGAGCACTTACTCAAGTCTGTCTTGCTCTACTCATTTGACACCTCAGTT GTTACTGCTCTAGAGCACTTACTCAAGTCTGTCTTGCTCTATTCCTTTGACACATCAGTT ***********************************
Chicken Turkey Ostrich	CAGGGACGTCAGAGCTGGGAAAGCGGGACCCCCAGGCTCCAGCCGGACTCCCGAGCCCTG CAGGGACGTCAGAGCTGGGAAAGCGGGACCCCCAGGCTCCAGCCGGACTCCCGAGCCCTG CAGGGACACCAGAATTGGGAAAGTGAGACTCCCAGGCTTCAGCCAGACTCCAGAGCCCTT ****** *** ******* **** **********
Chicken Turkey Ostrich	AGGTACATGAAGAGGCTGTATAAGATGTACGCCACCAAGGAGGGAATCCCAAAGGCCCAT CGGTACATGAAGAGGCTGTATAAGATGTATGCCACCAAGGAGGGAATCCCGAAGGCCCAC AGGTACATGAAGAGGCTATATAAGATGTCTGCCACCAAGGAGGGAATCCCAAAGGCCAAC **************************
Chicken Turkey Ostrich	AAGAGCCACCTCTATAACACTGTTCGACTTTTCACCCCGTGTTCTGAGTGCCAGCACCGC AAGAGCCACCTCTATAACACCGTTCGACTTTTCACCCCGTGTTCCGAGTGCCAGCACCGC AAAAGTCACCTTTATAACACTGTTCGACTTTTCACTCCGTGTTCTGAATGCAAGCACCGC ** ** ***** ******* ***************

Chicken Turkey Ostrich	CCCACTTCTTCCTTTACGTGCACGTGCCATTTGTCTGTGAAGGAGCATGATTTTTCT CCCACTTCTTCCTTTATGTGCACGTGCCATTTGTCTGTTAAGGAGCATGATTTTTCT CCCATTTCTTCTGCCACTACATGCATGTGCCATTTATCTGTTACGGAATATGATTTTTCT **** *****
Chicken Turkey Ostrich	AGCCAAGTATGTCCCAGCGTTTCACACTCCGTAGCTTTTAGCCTGCACTTTGAAGTTAGA AGCCAAGTATGTCCCAGCGTTTCGCACTCCGTAGCTTTTAGCCTGCACTTTGAAGTTAGA AGCCAAGTGTGTCCAAGTGTTTCACACTCTATAGCTCTTAGTTTGCACGTTGAAGTTAGA ******* **** ***** ***** ***********
Chicken Turkey Ostrich	AAGCGCAAGTGGGTTGAGATTGATGTGACTTCTTTTCTGCGGCCTCTCATTGCTACTAAC AAGCGCAAGTGGGTTGAGATCGATGTGACTTCTTTTCTGCAGCCTCTCATTGCTGCTAAC AAACGCAAGTGGGTTGAGATTGATGTGACTTCTTTTCTCCAGCCTCTAATTGCTACTAAC ** *********************************
Chicken Turkey Ostrich	AGGAGGAATATTCATATGGCTGTGAACTTCACTTGTCTGACGGGTAACCCGCAACATAAC AGGAGGAATATTCACATGGCTGTGAACTTCACTTGTCTTTTGGGTAACCCACAACATAAC AGGAGGAATATTCATATGGCTGTAAACTTCACTTGTCTAATGGGTGATCCACAACATAAC *************************
Chicken Turkey Ostrich	ACTAAACAGGATAATCTCATTAATGTGGCTCTGGTCCCCCCTTCTCTTCTTCTTTACCTA GCTAAACAGGATAATCTCATTAATATGGCTCTGGTCCCCCCTTCTCTTCTTCTTTACCTA ACTAAACAGGAAAATTCCATTAACATGGCACTGGTTCCCCCTTCTCTTCTTCTTTACCTG ********* *** ***** **** ************
Chicken Turkey Ostrich	AATGATACCAGCGAGCAAGCTTATCACAGGTGGAACTCACTTAGACACAGAAGGAAAAGC AATGATACCAGCGAGCAAGCTTATCACAGGTGGAACTCACTC
Chicken Turkey Ostrich	CCGGTGCGGCCCAAGCAAAGGAGCAGTCTGTTTGCTGATGTGACAGGTGATGAAGGAAG
Chicken Turkey Octrich	CAGAACACGCAGGGTAAAAGGGCATCTCGGCATCGAAGAGAAGAGAATCTGAAGGAAG
Chicken Turkey Ostrich	CCAGCGACTGCGCCTCAAAACTTGAGTGAATATTTCAAACAATTCCTGTTTCCTCAAAAC CCAGCAACTCCGCCTCAAAATTTGAGTGAATATTTCAAACAATTTCTGTTTCCTCAAAAC CCAGCAGCACCACCTTATAATTTGAGTGAATATTTCAAACAGTTTCTGTTTCCTCAGAAC ***** * * *** * ********************
Chicken Turkey Ostrich	GAGTGTGAGCTCCACAGCTTCCGTT, TAAGTTTTAGCCAATTAAAATGGGACAAATGGATA GAGTGTGAGCTTCACAGCTTCCGTTTAAGCTTTAGCCAACTAAAATGGGACAAATGGATA GAGTGTGAGCTTCACAACTTTCGTCTAAGTTTTAGCCAACTAAAATGGGACAAATGGATA ********** *** *** *** *** **********
Chicken Turkey Ostrich	ATAGCCCCGCATCGATACAGCCCTCCAGTACTGCAAGGGTGACTGTCCGAGGGTAGTCGG ATAGCACCGCATCGATACAGCCCTC-AGTACTGCAAGGGTGACTGTCCCAGGGTGGTTCG ATAGCACCTCATAGGTACAGCCCTC-AGTATTGCAAAGGTGACTGCCCAAGAGTAGTTGG ***** ** ** * ********* **** ********

gene in poultry species

chicken Turkey Ostrich	GCACCGTTATGGCTCTCCTGTACATACAATGGTGCAGAACATAATATACGAGAAGCTGGA GCACCGCTACGGCTCTCCTGTACATACAATGGTGCAGAACATAATATATGAGAAATTGGA CCATCGTTATGGCTCTCCCGTACATACAATGGTACAGAACATAATATATGAGAAACTGGA ** ** ** ******* *******************
Chicken Turkey Ostrich	CTCCTCTGTGCCCAGGCCCTCCTGTGTTCCTGCCGAATACAGCCCTCTGAGTGTCCTGAC CTCCTCTGTTCCCAGGCCCTCCTGTGTTCCTGCTGAATACAGCCCTCTGAGCGTCCTGAC CTCATCTGTTCCGAAGCC *** ***** ** ***
Chicken Turkey Ostrich	AATAGAGCCTGATGGCTCCATAGTCTACAAAGAATATGAGGACATGATAGCTACTAAGTG AATAGAGCCTGATGGCTCCATAGTCTACAAAGAATACGAGGACATGATAGCTACTAAGTG ANTAGAGCCCGACGGTTCTATAGTCTATAAAGAATATGAAGATATGATAGCTACTAAATG * ****** ** ** ** ******** ** ** ******
Chicken Turkey Ostrich	CACTTGTCGGTAGTGGGTGTACCCTAACCTGTCTTACACTTAGTTTGGGTGTTTTGC CACTTGTCGGTAGTGGGTGTATCCTAAGCTGCTTTACACTTAGTTTGGGTGTTTTTGCATT CACTTGTCGGTAG**********
Chicken Turkey Ostrich	TTTGCTAACAGCTCTTTCAAACTGAATAGGGTTTA
Chicken Turkey Ostrich	GCATGGAGTTAGTGGGTGTTTATATAACAGCACTGTTTTTGTATCCACTCAGCCTTTCTA GCATGGGATAAGAGGGTGTTTATATAATAGCACTATTTTTGTAGCCACTCAGTCTTTCTA
Chicken Turkey Ostrich	GAATGTGAAAACAAATGTAAATATCTTTTTTATTGGAATAATATTGCATTTGAACAGAAG GAACGTGAAAATAAATGTAAATATCTTTTT-ATTGGAATAATATTGCATTTGAACAGAAG
Chicken Turkey Ostrich	TGTTTTCAACTACCAGATGCCTCTTTTAAGCTGTTTTGTTACAGATTTAATAATAAATA
Chicken Turkey Ostrich	AGTGAATGCCTATCACACACATCACTTAAAAATGAATGCTTATCACTCATATCACTTAAATTCAGTGTCTGATGGTCTTCCTCT

Figure 1. Multiple sequence alignment of GDF9 DNA sequences of chicken, turkey and ostrich

Multiple alignment of protein sequences

Figure 2 shows g*rowth differentiation factor 9* protein sequences of chicken, turkey and ostrich which were aligned using CLUSTAL-W incorporated by Online Analysis Tools. The results showed that the percentage similarity of *GDF9* protein sequences of the selected poultry species ranged from 84-95% The percentage similarity of chicken and turkey protein sequences was found to be 94.74% and 83.63% between chicken and ostrich. On the other hand, turkey and ostrich shared 84,08% similarity in their *GDF9* protein sequences.

Chicken Turkey Ostrich	MGGTWRICVCFYCCLHWLSSSIQCSPRSRDRTAPDKVSGLLGAPELSELHPLLRLPKG MAGTWRICVCFYCCIQWLSSSIQCSPRSRDRAAPDKLSGLLEAPELSELHPLLRLPQG MGSPWRICVCFYCCIHWLSSSIQCSPRSRDHVVSDKTSGLLVAPEDYTSKLNPLLRLPKG **********::***********************
Chicken Turkey Ostrich	VSRGYALLPPLLEVLSDQGRQSWESGTPRLQPDSRALRYMKRLYKMYATKEGIPKAHKSH ASRGCALLPPLLKVLSDQGRQSWESGTPRLQPDSRALRYMKRLYKMYATKEGIPKAHKSH VRRGYALLPPLLKVLSDQGHQNWESETPRLQPDSRALRYMKRLYKMSATKEGIPKANKSH . ** ******: **************************
Chicken Turkey Ostrich	LYNTVRLFTPCSECQHRHRDPITGDFHSVDLLFNLDRVTALEHLLKSVLLYSFDTSVPTS LYNTVRLFTPCSECQHRHRDLITGDLHSVDLLFNLDRVTALEHLLKSVLLYSFDTSVPTS LYNTVRLFTPCSECKHRHRDLMKGDLHSVDLLFNLDRVTALEHLLKSVLLYSFDTSVPIS ************************************
Chicken Turkey Ostrich	-SFTCTCHLSVKEHDFSSQVCPSVSHSVAFSLHFEVRKRKWVEIDVTSFLRPLIATNRRN -SFMCTCHLSVKEHDFSSQVCPSVSHSVAFSLHFEVRKRKWVEIDVTSFLQPLIAANRRN SATTCMCHLSVTEYDFSSQVCPSVSHSIALSLHVEVRKRKWVEIDVTSFLQPLIATNRRN : * *****.*:***************************
Chicken Turkey Ostrich	IHMAVNFTCLTGNPQHNTKQDNLINVALVPPSLLLYLNDTSEQAYHRWNSLRHRRKSPVR IHMAVNFTCLLGNPQHNAKQDNLINMALVPPSLLLYLNDTSEQAYHRWNSLRHRRKSPVR IHMAVNFTCLMGDPQHNTKQENSINMALVPPSLLLYLNDTSEQAYHRWNSLRHRRKNPVW ******** *:***:** **:*****************
Chicken Turkey Ostrich	PKQRSSLFADVTGDEGRQNTQGKRASRHRREENLKEAPATAPQNLSEYFKQFLFPQNECE PKQRNSLFADVTGGQGRENTQGKRASRHRREENLKEAPATPPQNLSEYFKQFLFPQNECE PRQRNNLLADPMDGQGKENTQSKRASRRRRDEHLKEAPAAPPYNLSEYFKQFLFPQNECE *:***:**:*::***.******************
Chicken Turkey Ostrich	LHSFRLSFSQLKWDKWIIAPHRYSPQYCKGDCPRVVGHRYGSPVHTMVQNIIYEKLDSSV LHSFRLSFSQLKWDKWIIAPHRYSPQYCKGDCPRVVRHRYGSPVHTMVQNIIYEKLDSSV LHNFRLSFSQLKWDKWIIAPHRYSPQYCKGDCPRVVGHRYGSPVHTMVQNIIYEKLDSSV **.*********************************
Chicken Turkey Ostrich	PRPSCVPAEYSPLSVLTIEPDGSIVYKEYEDMIATKCTCR PRPSCVPAEYSPLSVLTIEPDGSIVYKEYEDMIATKCTCR PKPXEPDGSIVYKEYEDMIATKCTCR *:* ********************************

Figure 2. Multiple sequence alignment of GDF9 protein sequences of chicken, turkey and ostrich

Phylogenetic analysis

The evolutionary relationship among the DNA and protein sequences of *GDF9* gene of chicken, turkey and ostrich and other selected livestock species such as sheep, goat and cattle, are shown in Figure 3 and Figure 4, respectively. The relative taxa were grouped together in the bootstrap test (1000 replicates) and the percentage of the replicate trees is exhibited near the branches. The phylogenetic study revealed that the phylogenetic relationship of the chicken, turkey and ostrich grouped together with a bootstrap probability of 100%, equivalent to a confidence level of 99%. Chicken and turkey were found to share the last internal node on the phylogenetic trees

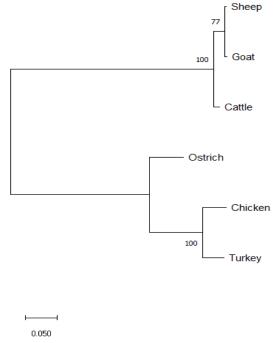


Figure 3. Phylogenetic tree showing the evolutionary relationship among poultry species and other selected livestock species using mRNA sequences.

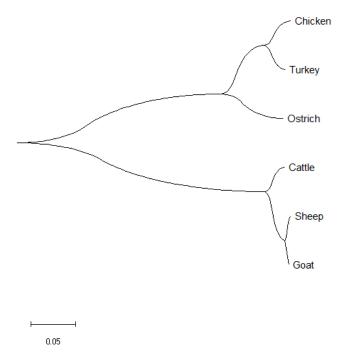


Figure 4. Phylogenetic tree showing the evolutionary relationship among poultry species and other selected livestock species using protein sequences.

Discussions

Bioinformatic tools, such as multiple sequence alignment and phylogenetic analysis, involve the use of computer softwares to make sense of great amounts of biological data (Madiajagan and Raj, 2019). The objective of the current study was to computationally investigate the molecular genetic variation of growth differentiation factor 9 gene in poultry species using multiple sequence alignment and phylogentic analysis whereby CLUSTAL-W was used to align DNA, and protein sequences and determine the percentage similarities while MEGA11 software was used to carry out the evolutionary study. Percentage similarity values between the GDF9 mRNA and Protein sequences of chicken, turkey, and ostrich ranged from 78.45-92.80% and 83.63-94.74%, respectively, meaning that the species are highly similar in function. These high levels of similarities in GDF9 gene sequences of chicken, turkey and ostrich is proof that the gene is very effective for its biological function improvement (Egom et al., 2019). The high similarity percentages also imply that the GDF9 gene at which each poultry species evolved is due to divergence, meaning that the species sharing similar ancestry acquired differences, gradually leading to speciation throughout the course of evolution (Gautam, 2020). The results are supported by Koonin and Galperin (2003), who documented that the likelihood that two sequences began independently of one another and became similar purely by coincidence decreases with increasing similarity between them. The phylogenetic studies revealed that since the selected poultry species clustered together into the same taxon, they have evolved from a common ancestor (Egom et al., 2019). The results further showed that chicken and turkey share a most recent common ancestor, as depicted by the last internal node which is furthest from the root of the evolutionary tree.

According to Egom *et al.* (2019), species sharing a most recent common ancestor will have a high percentage similarity value. This scenario is observed in the findings of the phylogenetic analysis of protein sequences in this study, whereby chicken and turkey share the highest similarity value and the most recent common ancestor. The reports, however, are in contradiction with the phylogenetic analysis of the DNA sequences which shows that chicken and ostrich do not share a most recent common ancestor regardless of having high percentage similarity of their *GDF9* gene sequences. Egom *et al.* (2019) and Divya *et al.* (2018) further reported that chicken and turkey do not share a most recent common ancestor thus in line with the phylogenetic analysis of the protein sequences and in contradiction with those of the DNA sequences. These differences may occur as a result of conducting the study with different genes. The study only used three poultry species; hence further studies can be done to improve egg production through *growth differentiation factor* 9 gene using other poultry species.

Conclusion

The study concluded that *GDF9* mRNA and protein sequences of chicken, ostrich and turkey were highly similar, and that the species were derived from a common ancestor thus these species can be improved similarly or through the other. The current study recommends that researchers focusing on *GDF9* for improving egg production in chicken might also improve egg production in ostrich and vice versa. It is also recommended that researchers may improve chicken and turkey for egg production under similar conditions.

Competing interest

There is no conflict of research interest.

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