Intermixing of Commercial Pure Breed Chickens with Indigenous (Sakini) Breed of Nepal

Neena Amatya Gorkhali¹, Chhiring Sherpa¹, Mana Raj Kolachhapatri², Bhoj Raj Pokharel¹, Nirajan Bhattachari² and Saroj Sapkota¹*¹

¹National Animal Breeding and Genetics Research Centre, Nepal Agricultural Research Council, Khumaltar, Lalitpur, Nepal; neenagorkhali@hotmail.com, SS: sarose.sapkota@gmail.com, ORCID: https://orcid.org/0000-0002-2930-2219
²Institute of Agriculture and Animal Science (IAAS), Rampur Chitwan, Nepal
³Agriculture and Forestry University, Rampur, Chitwan, Nepal NB: nbhattarai@afu.edu.np, ORCID: https://orcid.org/0000-0003-0325-6506

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ABSTRACT

The intermixing of pure breeds of commercial chickens with Nepalese indigenous flock of chickens has not been explored yet genetically. The study aims to investigate the status of interbreeding of Nepalese indigenous chickens with exotic breeds. Thirty three samples of indigenous chicken, Sakini, from three different ecological regions, Terai, mid-hill and high hill, were taken for the study. The 522 bp hyper-variable region of D loop mtDNA of each sampled population was PCR amplified and sequenced. The Neighbor joining tree revealed that most of Nepalese Sakini and commercial chickens such as New Hampshire, Rhode Island Red, White Leghorn, White Plymouth Rock and Barred Plymouth Rock lied in same clade. Our study provides first direct evidence that the Sakini was already bred with commercial breeds for the sake of enhancing production performance and also alarmed the government to introduce proper strategies and policies to avoid such genetic erosion.

Keywords: Sakini, mtDNA, intermixing, commercial breeds, genetic erosion.

सारांश

नेपाली स्थानीय जातका कुबु नंगे व्यवसायिक कुबु विच अन्तरजातिक प्रजननको विविध आनुभाषिक हिसाबले ज्ञात भएको हो तथा अनुसंधान भएको थिए। यस अध्ययनको मुख्य उद्देश्य विदेशी जात र नेपाली रैथाने जातका कुबु रातो बिचको मार्गर प्रजनन प्रक्रमको बारेमा अनुसंधान गर्नु र प्रयास छ। यस अध्ययनलाई गरेका प्रक्षेत्रहरू जातका साकिनी कुबु रातो रातको नमुनाहरू विभिन्न तिन अन्तर अन्तर व्यवसायिक ज्ञेय र न्याय र उद्योग प्राधिकारीको विभिन्न तिन अन्तर व्यवसायिक ज्ञेय र न्याय र उद्योग प्राधिकारीको विभिन्न तिन अन्तर

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INTRODUCTION

The indigenous breeds, which serve as a major gene pool, are mainly raised in most of the developing countries (Bett et al., 2011). In Nepal, the poultry farming of both commercial and native stocks are in increasing trend. Dhakal (2005) revealed that the poultry sector solely encompasses 4% of the agricultural GDP in Nepal. Neokane and Gorkhali (2008) reported that 50% of total poultry population consisted of native breeds in Nepal. Some of the documented indigenous chicken breeds of Nepal are Sakini (Normal feather), Ghatikhule (naked neck) and Pwakh ulte (frizzled feather) (Neokane and Gorkhali, 2008). Among three indigenous chicken breeds, Sakini shares more than 50 percentage in total national poultry population (Country Report, 2014).

Sharma (2012) claimed that the poultry farming of commercial chickens have proliferated mainly in urban areas where as indigenous chickens are mostly limited in rural areas of the country because of poor access of roads and transportations. Even though these native chickens are primarily used in Family Poultry Production System (FPPS) in Nepal, the productive performances of these chickens are quite low (Sapkota et al., 2020). The productive performance of native chickens is relatively lower than other commercial breeds but they are embodied with the major genes which aid them to adapt harsh environmental conditions and resist against many diseases and parasites (Bhatti et al., 1990).

Due to poor productivity in indigenous chickens with reference to the growth, egg size and the egg production, the farmers have been introducing the pure exotic breeds in their farming systems for crossbreeding with indigenous breeds (ABD 2018; Tadella et al., 2020). The introgression of the exotic breeds on the native stocks results in the genetic erosion. Subsequently, the loss of positive attributes in indigenous breeds may occur (Gorkhali et al., 2020; Sharma et al., 2001).

Among the indigenous chicken breeds of Nepal, Sakini is the principal breed, but it has not been adequately characterized molecularly (Gorkhali et al., 2013). Sapkota et al. (2020) reported that the red jungle fowl from India was the progenitor of Sakini but the introgression of commercial chicken into this principal breed has not been studied in detail. Thus, this study aims to assess the status of interbreeding of principal breed, Sakini, with other commercial breeds.

MATERIALS AND METHODS

Sampling

Thirty three blood samples of Sakini chickens were taken from three different agro-ecological regions of Nepal (Terai, mid-hill and high hill). DNA extraction was done by following the protocol provided by Promega DNA extraction kit (Promega Korea, Republic of Korea). The quality of DNA was further assessed by using a Nanodrop spectrophotometer (Thermo Scientific™ NanoDrop 2000, China). The DNA samples were safely preserved at -20 degree Celsius until the test. Forty-two (42) reference sequences including commercial chickens, indigenous chickens and Red jungle fowl of different countries were used for analysis.

PCR amplification and sequencing

In order to amplify 522 bp long hyper-variable region of D loop mtDNA, the set of primer named L16750 and H547 (Niu et al., 2002) were used (Table 1). The final total volume of PCR reaction was made 25μl by adding 13 μl of PCR 2X Master Mix, 4 μl of molecular grade nuclease-free water, 5 μl of template DNA, and 1.5 μl of each forward and reverse primers. The optimum PCR condition was as follows: denaturation at 95°C for 1 minute, annealing at 63°C for 1 minute and extension at 72°C for 2 minutes. The amplification of 522 bp hyper-variable regions was checked by running on 1.5% agarose gel followed by visualizing under UV light by comparing with a DNA molecular weight marker (100 bp ladder). 10 μl of the PCR product of each sample which showed fine amplification on agarose gel was sent to the Department of Animal Genetic Resources, National Livestock Research Institute, Rural Development Administration, South Korea for sequencing. The sequenced data were then used for comparing with mtDNA sequences of commercial chicken breeds and indigenous breeds from different countries retrieved from National Center for Biotechnology Information (NCBI).
Data analysis
The obtained 33 mtDNA sequences of Sakini were edited by using CHROMAS software and aligned against 42 reference sequences of commercial chickens, RJF and domestic chicken of different countries by using ClustalW program of MEGA 5 software (Thompson et al., 1994). By following Kimura-2-parameter model of MEGA 5 software, Neighbour joining (NJ) phylogenetic tree was constructed (Tamura et al., 2011).

Table 1: The oligonucleotide sequences of L16750 and H547.

| S.N | Primer | Oligonucleotide                  |
|-----|--------|---------------------------------|
| 1   | L16750 | 5’-AGG ACT ACG GCT TGA AAA GC-3’ |
| 2   | H547   | 5’-ATG TGC CTG ACC GAG GAA CCA G-3’ |

RESULTS
The Neighbour joining (NJ) tree revealed that the Nepalese Sakini chickens shared the same cluster with various commercial chicken breeds such as New Hampshire, White Plymouth Rock, Barred Plymouth Rock, Rhode Island Red and White Leghorn. Also major of these Sakini population share the same mtDNA D loop of domestic breeds from China and India.

DISCUSSION
The commercial pure breeds such as New Hampshire was imported from India to Nepal in 1964 by the help of US funded project (FAO, 2014). Many farmers have prioritized commercial breeds as a poultry farming since Nepalese indigenous chicken breeds have lower production performance in terms of growth, egg production and egg size (ABD, 2018). Assessing the financial benefits of commercial
poultry farming, Government of Nepal initiated various approaches and programmes to distribute it to Nepalese farmers aiming poverty reduction (Kayastha, 2010). Paucity of information is found in terms of Nepalese poultry conditions; it makes arduous to estimate the time period from when the actual crossbreeding between commercial and indigenous chicken occurred (FAO, 2014). It is reported that crossbreeding of indigenous chickens with the pure breeds of commercial chickens is popular to enhance genetic potentiality as a result an increase in production performances can be attained (Tadelle et al., 2020). Nepalese farmers may have undergone same procedure for improving the production performance of Nepalese native chickens (ABD, 2018). Yet, no promising evidence has been provided which reveals that there has been intermixing of commercial breeds of chickens with indigenous breed.

Our result provides the first evidence that our Nepalese breed, *Sakini*, was already bred with exotic breeds such as New Hampshire, white Plymouth Rock, Barred Plymouth Rock, Rhode Island Red and White Leghorn. Our study claiming *Sakini* to be close with domestic chicken breeds of India support the findings made by Bhurtel (2011) and Sapkota et al (2020). Such intermixing results in genetic erosion; as a result the native breeds may lose genes of positive attributes which are helping them to combat against extreme climatic conditions and diseases (Hosny 2005). If such intermixing continues for long run, then the Nepalese chicken will lose its genetic diversity; as a result they may no longer be able to resist against several diseases and cannot be habitant of extreme climates (Sharma et al., 2001). Our study draws attention to the government of Nepal for formulating judicious crossbreeding strategies so that the positive attributes of the indigenous chickens will not get lost. As a result genetic diversity among *Sakini* population can pass from generation to generation without loss of favourable genes.

CONCLUSION

This study reports the first evidence that the Nepalese *Sakini* breeds was already bred with different commercial pure breeds of chickens for improving genetic potentiality to enhance the productivity of the native chickens. Defined government policies with regard to crossbreeding commercial chicken with indigenous chicken populations must be formulated to prevent genetic erosion.

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