Cloning of the short-tailed Gyeongju Donggyeong dog *via* SCNT: conserving phenotypic inheritance

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ABSTRACT. Somatic cell nuclear transfer is a useful tool to maintain genetic information of animals. The Gyeongju Donggyeong dog is a breed registered as natural monument in Korea. The unique feature of the Donggyeong dog is its tail, as the Donggyeong dog can be classified as either short tailed or tailless. The aim of this study was to preserve the Donggyeong dog's unique feature by cloning. Fibroblasts were obtained from a short-tailed Donggyeong dog. *In vivo* matured oocytes were enucleated, microinjected with a donor cell and fused electrically. Reconstructed embryos were transferred to six recipient dogs. One surrogate became pregnant, and one short-tailed Donggyeong dog was delivered. This study demonstrated that the phenotype of the Donggyeong dog could be conserved by somatic cell nuclear transfer. KEY WORDS: Gyeongju Donggyeong dog, phenotype, somatic cell nuclear transfer

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Since the production of the first cloned dog, Snuppy (Afghan hound) [8], several species, such as the beagle [3], toy poodle [4], retriever [9], border collie [7] and Pekingese [12], have been cloned by somatic cell nuclear transfer (SCNT). Among the many breeds that need to be saved from extinction, the Sapsaree, one of the Korean natural monument dogs, has been produced by SCNT [5]. The Gyeongju Donggyeong dog has been considered a natural monument since 2012 (Cultural Heritage Administration of Korea, number: 540). The name Gyeongju Donggyeong dog originated from the capital of the ancient Silla kingdom in Korea. The Donggyeong dog has the oldest history among the Korean natural monument breeds, and it is referred to in many historic documents, such as Dongkyung jabki (published in AD 1669) and Sungho sasul (published in AD 1740). Despite the Donggyeong dog's high historical value, only about two hundred individuals remain in Gyeongju, and it is classified as endangered [1, 2, 10]. It is essential to save such a valuable breed from extinction and maintain a pure descent. Accordingly, the aim of this study was to clone the Donggyeong dog by SCNT and observe the similarity of phenotypes between the cloned and cell donor dogs.

In this study, mixed-breed dogs between one to five years were used as oocyte donors and embryo recipients. The study was conducted in accordance with recommendations described in "The Guide for the Care and Use of Laboratory

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For preparation of donor cells, skin tissue was isolated by an aseptic surgical method from a three-month-old female Donggyeong dog (Fig. 1A). Recovery of *in vivo* matured oocytes was performed from oviducts approximately 72 hr after ovulation. Prediction of ovulation, preparation of matured oocytes, the process of SCNT and the transfer method for cloned embryos were described previously [6,



Fig. 1. Pictures of the cell donor and cloned Donggyeong dogs.A) cell donor dog at three months old. B) cloned dog at 1 day after birth. C) tail length of cell donor dog. D) tail length of cloned dog.

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Recipient	No. oocytes donors	Oocyte status ^{a)}	No. reconstructed couplets	Pregnancy	No. cloned dogs
А	2	Mature	15	_	_
В	1	Mature	6	_	-
С	3	Mature and early aged	21	+	1
D	1	Mature	9	_	_
Е	3	Early mature, mature, immature	29	_	_
F	2	Early mature	18	-	_
Total (n=6)	12		98	1 (16.6%) ^{b)}	1 (0.01%) ^{c)}

Table 1. In vivo development of cloned embryos by SCNT using somatic cells derived from donor dog

a) Status of *in vivo* oocytes flushed from oviducts approximately 72 hr after ovulation. b) The percentage is based on the total number of recipient dogs. c) The percentage is based on the total number of transferred embryos.

8]. Reconstructed embryos (n=98) were transferred into oviducts of six recipient dogs that were naturally synchronized. One recipient dog was confirmed to be pregnant by ultrasonography 26 days after embryo transfer (pregnancy rate: 16.6%) (Table 1). Pregnancy was maintained to term, and one healthy female Donggyeong dog weighing 320 g was delivered by cesarean section 59 days after embryo transfer (Fig. 1B).

In order to identify the origin of the mitochondrial DNA (mtDNA) in the cloned dog, the genomic DNA was used for canine mtDNA (GenBank accession no U96639 v.2 and 650 bases) analysis. Based on the results, we identified that the cloned dog had identical mtDNA sequences to those of the domestic oocyte donor and surrogate dog (Table 2). To clone the Donggyeong dog, mixed-breed dogs were used as oocyte donors and recipients. Since the oocyte donor dog and recipient dog showed the same information for mtDNA in the mtDNA analysis, it is difficult to distinguish which dog's mtDNA was transferred to the cloned dog (Table 2). Many studies about dog cloning proved that the cloned dog's mtDNA was transferred solely from the oocyte donor [4, 5, 7]. Similarly, in the present study, the mtDNA of the cloned dog might have been transferred from the oocyte donor dog. In addition, to determine parentage, DNA extraction and microsatellite analysis with canine-specific markers were performed following the protocol of our previous study [6]. The results of the parentage analysis indicated that the cloned Donggyeong dog was genetically identical to the cell donor Donggyeong dog (Table 3).

It has been reported that a cloned toy poodle had the same coat color as the somatic cell donor dog [4] and that beagles cloned from fetal fibroblasts had similar coat spotting [3]. In the present study, the cloned Donggyeong dog also had a phenotype similar to that of the cell donor dog. The unique feature of the Donggyeong dog is that it can be classified as natural short tailed or tailless. According to information from the Korean Gyeongju Donggyeong Dog Association, the short-tailed Donggyeong dog over twelve months old has a tail length of around 11.38 \pm 2.44 cm and five to nine of coccygeal vertebrae in radiographic observations [2, 10]. The tailless Donggyeong dog has one to four of coccygeal vertebrae, and adult dogs (over twelve months old) have a tail length of around 6.3 ± 2.81 cm. In order to identify the number of caudal vertebral bodies, the dorsal radiographic

views of the caudal vertebral column in the cloned dog and donor dog were compared. The number of coccygeal vertebral bodies was counted from the sacrum of the dorsal surface of the vertebral body. Based on the dorsal radiographic view, the cell donor dog had six coccygeal vertebral bodies, while the cloned dog had seven coccygeal vertebral bodies (Fig. 2A and 2B). Even though they had different numbers of coccygeal vertebral bodies, they could both be categorized as short-tailed Donggyeong dogs. It has been reported that despite cloned dogs having the same genetic information, they can have different dental development; one cloned dog had normal dental formulas, another cloned dog was missing one permanent molar tooth on the left side, and another cloned dog was missing one permanent molar tooth on the right side [11]. We cannot explain the reason for these differences between a donor dog and a cloned dog, but they might be associated with epigenetic modification during the cloning procedure. Therefore, further studies should be encouraged to analyze the epigenetic effects on the caudal vertebra features of cloned dogs.

A Donggyeong dog, which is considered an endangered breed and needs to be saved from extinction, was clones using SCNT for the first time. Furthermore, the cloned Donggyeong dog could be classified as short tailed, which is the same as the cell donor Donggyeong dog. The current study demonstrated that SCNT could not only be used for conserving a specific breed of dog but also that it could ensure inheritance of a unique phenotypic feature of a dog breed. To determine the relationship between the coccygeal vertebrae and epigenetic modification, further studies need to analyze epigenetic mechanisms, which can influence coccygeal vertebra development.

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	Nucleotide positions ^{a)}																		
Sample ID	15518	15526	15595	15612	15627	15632	15639	15643	15652	15665	15800	15814	15815	15912	15955	16003	16025	16083	16105
Reference	Α	С	С	Т	Α	С	Т	Α	G	Т	Т	С	Т	С	С	Α	Т	Α	Т
Cell donor	Α	С	С	Т	А	С	А	А	G	Т	Т	Т	Т	С	С	А	С	А	Т
Cloned dog	Α	С	С	Т	G	С	А	А	А	С	Т	Т	Т	С	С	А	Т	А	Т
Surrogate	Α	С	С	Т	G	С	А	А	А	С	Т	Т	Т	С	С	А	Т	А	Т
Oocyte donor 1	Α	С	С	Т	G	С	А	А	А	С	Т	Т	Т	С	С	А	Т	Α	Т
Oocyte donor 2	Α	С	С	Т	G	С	А	А	А	С	Т	Т	Т	С	С	А	Т	Α	G
Oocyte donor 3	C	Т	Т	С	Α	Т	G	G	Α	Т	С	Т	С	Т	Т	G	Т	G	Т

Table 2. Sequence alignments within 628 bases of the hypervariable region of mitochondrial DNA

a) The nucleotide positions were numbered according to GenBank accession no. U96639 v.2, and 650 bases (from 15461 to 16110) were examined.

 Table 3.
 Microsatellite genotyping of cell donor, cloned, surrogate and oocytes donor dogs using specific canine DNA markers

NAME	Cell donor	Cloned	Surrogate	Oocyte donor 1
PEZ2	130 / 130	130 / 130	126 / 122	126 / 126
PEZ10	298 / 282	298 / 282	282 / 282	282 / 262
PEZ16	298 / 290	298 / 290	302 / 286	302 / 282
CPH4	149 / 137	149 / 137	141 / 141	141 / 141
PEZ17	222 / 214	222 / 214	218 / 202	210 / 210
CPH12	207 / 207	207 / 207	203 / 193	193 / 193



- Fig. 2. Comparison of the number of coccygeal vertebral bodies of the cloned Donggyeong dog (20 days after birth) and a donor Donggyeong dog (six months old) using digital radiographic views. A) a dorsal radiographic view of a portion of the caudal vertebral column of a cell donor dog is shown to illustrate measurements obtained for the sacrum (white bracket) through to the last coccygeal vertebra. B) dorsal radiographic view of the cloned dog. The coccygeal vertebral number was measured as the number from the dorsal surface of the sacrum. The cell donor dog had six coccygeal vertebral bodies, whereas the cloned dog had seven coccygeal vertebral bodies.
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