The development and application of genome editing technology in ruminants: a review

Mengke YUAN1,2*, Yuanpeng GAO1,2*, Jing HAN1,2, Teng WU1,2, Jingcheng ZHANG1,2, Yongke WEI1,2, Yong ZHANG (✉)1,2

1 College of Veterinary Medicine, Northwest A&F University, Yangling 712100, China
2 Key Laboratory of Animal Biotechnology (Ministry of Agriculture), Northwest A&F University, Yangling 712100, China

Abstract Transgenic ruminants are a valuable resource for both animal breeding and biomedical research. The development of transgenic breeding is proceeding slowly, because it suffers from low efficiency of gene transfer and possible safety problems from uncontrolled random integration. However, new breeding methods combined with genome editing and somatic cell nuclear transfer or microinjection can offer an economic and efficient way to produce gene-edited ruminants, which can serve as bioreactors or have improved disease resistance, animal welfare and product quality. Recent advances in precise genome editing technologies, especially clustered regularly interspaced short palindromic repeat (CRISPR)/Cas9 nucleases, are enabling the systematic development of gene-edited ruminant production. This review covers the development of gene-edited ruminants, the particulars of site-specific engineered nucleases and the state of the art and new insights into practical applications and social acceptance of genome editing technology in ruminants. It is concluded that the production of gene-edited ruminants is feasible and through improvements in genome editing technology it is possible to help feed the world.

Keywords bioreactors, breeding, engineered endonucleases, genome editing, ruminants

1 Introduction

Gene-edited ruminants have a potentially broad range of applications in the improvement of product quality and animal welfare, the enhancement of disease resistance and the production of biomedical materials[1-4]. Previously, transgenic ruminants were obtained by microinjection transferring the exogenous DNA into the cytoplasm of zygotes[5]. However, suffering from high rate of embryo chimerism and low gene-edited animal birth rates, microinjection was and remains inefficient for gene targeting, especially in monotocous species, such as cattle[6,7].

For rapidly-reproducing polytocous animals, such as mice, many progenies stably expressing the intended gene can be selected from the offspring generated by micro-injection of pluripotent embryonic stem cells (ESCs) containing the exogenous gene into blastocysts and transferring the embryos to foster mothers[8]. While for ESC lacking ruminants (goats[9], cattle[10,11] and sheep[12,13]), somatic cell nuclear transfer (SCNT) technology is available. SCNT based cloning technology transfers a somatic cell known to carry and/or silence the gene of interest to an enucleated oocyte, and then through electrofusion and chemical activation to obtain reconstructed embryos with a predictable offspring genotype and reliable efficiency[14].

The latest genome editing technology exploited as a powerful tool for identifying gene function and curing genetic diseases[15] relies on site-specific engineered endonucleases (EENs), including zinc finger nucleases (ZFNs), transcription activator-like effector nucleases (TALENs), and clustered regularly interspaced short palindromic repeat (CRISPR)/Cas9 nucleases, triggering efficient DNA double-strand breaks (DSBs) at target sites[16] and DSBs mediated gene knockin (KI), knockout (KO), or substitution via intracellular DNA damage repair pathways[17]. In recent years, type II bacterial CRISPR/Cas9 has come to represent the most efficient tool for the further advancement of gene targeting strategies due to the high efficiency and simplicity of targeting any locus for cleavage with a single protein and a programmable single-guide RNA (sgRNA).

Here we review research progress in genome editing in...
ruminants and the structure and characteristics of EENs and provide an overview in detail of the application of genome editing technology to inspire further solutions for efficient and precise gene modification in ruminants.

## 2 Research progress in genome editing in ruminants

Dolly, the first sheep somatically cloned via SCNT, was born in 1996 in Scotland and demonstrated that mammals could be cloned from adult cells, especially cultivated somatic cells containing the intended gene modification\[^{18}\]. This showed that researchers can obtain gene-edited ruminants with predictable offspring genotype and reliable efficiency\[^{19}\]. The first gene-edited ruminant offspring was produced via SCNT in 2000\[^{20}\]. A common strategy for the production of gene-edited ruminants was to deliver sufficient donors, containing the intended gene or a DNA fragment flanked by homologous arms, as templates into the intended cells. Intracellular spontaneously created DSBs are principally repaired by the non-homologous end-joining (NHEJ) pathway and the competing homologous recombination (HR) pathway\[^{21}\]. Therefore, the intended gene or the DNA fragment could be integrated into the genome at DSB sites by the HR pathway to achieve target gene KO or KO\[^{16}\]. However, this method was limited in the production of gene-edited ruminants, due to the low frequency of DSBs and HR repair.

The next major advance was the advent of site-specific EENs, including ZFNs, TALENs and CRISPR/Cas9, which enabled direct targeting of a sequence for mutation, KO or site-specific KI in the genome\[^{16}\]. The first ruminants produced using EENs were cattle with a β-lactoglobulin (BLG) gene modified via ZFNs in 2011\[^{22}\]. During the process of NHEJ repair, small inserts and/or deletions (indels) are introduced at the DSB sites and may result in frameshifts and corresponding functional gene KO\[^{16}\]. Gene-edited ruminants with biallelic KO of the intended gene were obtained, combined with SCNT technology, owing to the high frequency of NHEJ repair\[^{23,24}\]. Also, researchers utilized microinjection by transferring the EENs mRNA or plasmids into the cytoplasm of zygotes, to produce individuals with the intended gene KO\[^{25,26}\].

The ability of homology-directed repair (HDR)-based gene editing to precisely integrate interested genes of the same or different species into the genome of targeted animal endows scientists with a powerful tool to directionally and quickly generate gene-edited varieties free from the limitations of painstaking traditional breeding methods, especially for disease resistance breeding\[^{1,7,27}\]. Through the introduction of DSBs at target sites, site-specific EENs enormously stimulate HR\[^{28}\] and significantly improve HDR-mediated gene KI efficiency, which is the key to produce gene-edited ruminants lacking ESC and induced pluripotent stem cell\[^{14,29}\].

Gene-edited ruminants have been obtained mainly by these technologies. For gene-edited cows, the main production steps (Fig. 1) and the main advantages and disadvantages of SCNT and microinjection technology are presented in Table 1.

## 3 Structure and characteristics of EENs

### 3.1 ZFNs and TALENs

Derived from eukaryotic transcription factors, ZFNs and TALENs are composed of one FokI nuclease domain and individual DNA recognition units assembled DNA binding domain\[^{30}\]. Every ZFN unit (about 30 amino acids)
combines with a single zinc atom and recognizes a 3-bp DNA sequence\cite{31}, while a TALE unit is composed of engineered TALE repeat arrays (33–35 amino acids) and recognizes a nucleotide, namely, NN, NI, HD and NG, recognizing guanine, adenine, cytosine and thymine, respectively\cite{32}. Once ZFNs or TALENs bind to the target sites, the dimerized FokI nuclease domains forcefully induce two nicks 5- or 6-bp apart on the individual strands of DNA to form a precise DSB at the target site\cite{33}, and then the endocellular DNA damage repair pathway could be triggered to KO endogenous functional genes\cite{33} or KI exogenous genes\cite{35}. However, designing, constructing and screening efficient ZFNs and TALENs is expensive and time consuming for most molecular biology laboratories\cite{36} and has relatively low efficiency\cite{37,38}. Serious off-target effects\cite{39–41} of ZFNs and TALENs were also reported in a few studies, which limited their development and application in the production of gene-edited ruminants.

### 3.2 CRISPR/Cas9

Unlike the other technologies, the type II CRISPR/Cas9 system has been most widely exploited for gene editing due to the simple requirements for a crRNA, tracrRNA and a Cas9 protein for successful in vitro DNA cleavage\cite{42,43}. In addition, the crRNA and tracrRNA can be artificially fused as a sgRNA to further simplify this system\cite{44}. The natural crystal structure of *Streptococcus pyogenes* Cas9 (SpCas9)-sgRNA-DNA complex indicate that the sgRNA-DNA heteroduplex divides the Cas9 protein into two lobes (a recognition lobe and a nuclease lobe) connected by a bridge helix\cite{45}. The recognition lobe comprises Rec1 and Rec2 domains, while the nuclease lobe includes RuvC, HNH and PID domains. The recognition of target DNA is initiated through Cas9 protein binding to the protoscaler adjacent motif (PAM) sequence, but the sgRNA-DNA complex is crucial for the progressive recognition process. Once the guide sequence in sgRNA is complementary to the target DNA, the RuvC and HNH nuclease domains will be allosterically activated to cleave the target DNA at a site three base-pairs downstream of PAM and subsequently produce a DSB\cite{15,44}. The CRISPR/Cas9 system is cheap, highly efficient and easy to use even for novices, and its cleavage site can be as accurate as a single base\cite{44}. Therefore, since the first CRISPR/Cas9 mediated gene editing in mammalian cells\cite{46,47}, this well-established system has been a widely utilized platform for gene-edited ruminant production\cite{1,48}.

### 4 Applications of genome editing technology in ruminants

Traditional selective breeding and crossbreeding are time consuming\cite{4} and only improve existing production traits, whereas gene-edited ruminants breeding offers particularly attractive possibilities for generating characteristics that do not exist in nature. The development of EEN-mediated genome editing technology was essential for increasing the efficiency of gene targeting, which promises to greatly improve the traits of disease resistance, animal welfare and product quality in ruminants, and increase the safety, control and environmental benefits of gene-edited animal breeding. Gene-edited ruminants using EENs also have been widely adopted as bioreactors, thereby greatly contributing to the improvement of human health and alleviation of disease (Table 2).

<table>
<thead>
<tr>
<th>Technology</th>
<th>Advantages</th>
<th>Disadvantages</th>
</tr>
</thead>
<tbody>
<tr>
<td>SCNT</td>
<td>Predictable offspring genotype</td>
<td>Higher rate of midwifery</td>
</tr>
<tr>
<td></td>
<td>Consistent expression of target gene</td>
<td>Lower production efficiency* (1%–20%)</td>
</tr>
<tr>
<td></td>
<td>100% gene editing efficiency</td>
<td>Not induced at term</td>
</tr>
<tr>
<td></td>
<td>Saving time and costs</td>
<td>Increased developmental defects</td>
</tr>
<tr>
<td></td>
<td>Allows selection of gender and elite genetic background</td>
<td>Greater risk of recessive genetic diseases</td>
</tr>
<tr>
<td>Microinjection</td>
<td>Relatively lower rate of midwifery</td>
<td>Chimerism</td>
</tr>
<tr>
<td></td>
<td>15%–40% higher production efficiency</td>
<td>Not particularly useful for insertion of exogenous gene</td>
</tr>
<tr>
<td></td>
<td>Easy to perform micromanipulation</td>
<td>Low number of zygotes</td>
</tr>
</tbody>
</table>

Note: *Pregnancy rate is low but the rate of abortion is high.

Table 1: Advantages and disadvantages of somatic cell nuclear transfer (SCNT) and microinjection technology for gene-edited cows

<table>
<thead>
<tr>
<th>Technology</th>
<th>Advantages</th>
<th>Disadvantages</th>
</tr>
</thead>
<tbody>
<tr>
<td>SCNT</td>
<td>Predictable offspring genotype</td>
<td>Higher rate of midwifery</td>
</tr>
<tr>
<td></td>
<td>Consistent expression of target gene</td>
<td>Lower production efficiency* (1%–20%)</td>
</tr>
<tr>
<td></td>
<td>100% gene editing efficiency</td>
<td>Not induced at term</td>
</tr>
<tr>
<td></td>
<td>Saving time and costs</td>
<td>Increased developmental defects</td>
</tr>
<tr>
<td></td>
<td>Allows selection of gender and elite genetic background</td>
<td>Greater risk of recessive genetic diseases</td>
</tr>
<tr>
<td>Microinjection</td>
<td>Relatively lower rate of midwifery</td>
<td>Chimerism</td>
</tr>
<tr>
<td></td>
<td>15%–40% higher production efficiency</td>
<td>Not particularly useful for insertion of exogenous gene</td>
</tr>
<tr>
<td></td>
<td>Easy to perform micromanipulation</td>
<td>Low number of zygotes</td>
</tr>
</tbody>
</table>

Note: *Pregnancy rate is low but the rate of abortion is high.

Bioreactors based on gene-edited ruminants were developed for biomedical materials, such as enzymes and hormones\cite{54}. Ruminants have huge mammary glands and secrete large quantities of milk, with milking cows producing up to 8000 L·yr⁻¹ (cf.800 L·yr⁻¹ for goats and 500 L·yr⁻¹ for sheep). Due to the similarities in glycosylation patterns between the proteins secreted by the mammary gland tissue of ruminants and human native proteins\cite{54}, the development of technologies related to the purification of recombinant proteins from milk provides a

4.1 Bioreactors

Bioreactors...
mature and widely used system for recombinant proteins and makes ruminants most suitable species for recombinant proteins production\cite{54,55}. Researchers have introduced a therapeutic gene encoding human α1-antitrypsin (AAT) into α1(I) procollagen locus in primary ovine fetal fibroblasts by HR-mediated gene editing \cite{20}. Nuclear transfer with four independent donor cell lines resulted in the production of 14 gene-edited sheep, three of which lived for over one year. The analysis of hormonally induced milk of one of the three lambs showed the expression of AAT at a concentration of 650 mg mL\(^{-1}\). In contrast, previous research reported the highest level of AAT production observed in milk of gene-edited sheep produced by random integration only reached 18 mg mL\(^{-1}\).

AAT is available for the treatment of AAT deficiency in patients with clinically evident emphysema\cite{56}. Compared with random integration, gene-edited sheep produced by HR-mediated AAT gene integration showed a dramatic increase in yield.

Melatonin is a potent antioxidant widely used for medicinal purposes. In 2017, researchers report the first functional arylalkylamine N-acetyltransferase (AANAT) and acetylserotonin methyltransferase (ASMT) gene-edited animal model for high level of melatonin production in the milk of gene-edited sheep\cite{48}. Thirty-four gene-edited lambs were liveborn: seven contained AANAT, two contained ASMT, whereas 25 contained both AANAT and ASMT. CRISPR/Cas9 and microinjection technologies were combined to generate gene-edited sheep at a rate of up to approximately 35%, compared with 5% without EENs\cite{57}.

### 4.2 Disease resistance

Using genome editing technology, biomedical materials can be produced in ruminants. However, there are safety concerns about the products of ruminants infected with certain diseases, especially mastitis and prion diseases. Both diseases also affect animal health and welfare, causing substantial financial strain to the dairy industry worldwide.

Mastitis is one of the most costly diseases in the dairy industry and has a great influence on ruminants’ health and welfare. The industry suffers from high costs brought

<table>
<thead>
<tr>
<th>Application</th>
<th>Species</th>
<th>Target</th>
<th>Trait/Goal</th>
<th>Methods</th>
<th>Time</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bioreactors</td>
<td>Sheep</td>
<td>Insertion of <em>AAT</em></td>
<td>Treatment of α1-antitrypsin (AAT) deficiency</td>
<td>HR/SCNT</td>
<td>2000</td>
<td>\cite{20}</td>
</tr>
<tr>
<td>Sheep</td>
<td>Insertion of <em>AANAT</em> and <em>ASMT</em></td>
<td>Melatonin-enriched milk</td>
<td>CRISPR/Cas9 and zygote injection</td>
<td>2017</td>
<td>\cite{48}</td>
<td></td>
</tr>
<tr>
<td>Disease resistance</td>
<td>Cows</td>
<td>Insertion of lysostaphin</td>
<td>Lysostaphin milk</td>
<td>ZFN Nickase/SCNT</td>
<td>2013</td>
<td>\cite{27}</td>
</tr>
<tr>
<td>Cattle</td>
<td>KO of PrP</td>
<td>Lacking prion protein</td>
<td>HR/SCNT</td>
<td>2007</td>
<td>\cite{49}</td>
<td></td>
</tr>
<tr>
<td>Sheep</td>
<td>KO of PrP</td>
<td>Lacking prion protein</td>
<td>HR/SCNT</td>
<td>2001</td>
<td>\cite{50}</td>
<td></td>
</tr>
<tr>
<td>Cattle</td>
<td>Substitution of CD18</td>
<td>Abolish cytolysis of leukocytes</td>
<td>ZFNs/SCNT</td>
<td>2016</td>
<td>\cite{51}</td>
<td></td>
</tr>
<tr>
<td>Cattle</td>
<td>Insertion of <em>NRAMP1</em></td>
<td>Tuberculosis-resistant</td>
<td>Cas9 nickase/SCNT</td>
<td>2017</td>
<td>\cite{1}</td>
<td></td>
</tr>
<tr>
<td>Cattle</td>
<td>Insertion of <em>SP110</em></td>
<td>Tuberculosis-resistant</td>
<td>TALE nickase/SCNT</td>
<td>2015</td>
<td>\cite{7}</td>
<td></td>
</tr>
<tr>
<td>Animal welfare</td>
<td>Cattle</td>
<td>Integression of POLLED allele</td>
<td>Polled dairy cattle</td>
<td>TALEN/SCNT</td>
<td>2016</td>
<td>\cite{3}</td>
</tr>
<tr>
<td>Milk</td>
<td>Cattle</td>
<td><em>BLG</em> biallelic mutations</td>
<td>Modification of <em>BLG</em></td>
<td>ZFNs/SCNT</td>
<td>2011</td>
<td>\cite{22}</td>
</tr>
<tr>
<td>Cows</td>
<td><em>BLG</em> biallelic KO</td>
<td><em>BLG</em>-free milk</td>
<td>ZFNs/SCNT</td>
<td>2018</td>
<td>\cite{23}</td>
<td></td>
</tr>
<tr>
<td>Cattle</td>
<td><em>BLG</em> KO</td>
<td>Milk free of mature <em>BLG</em></td>
<td>TALEN and zygote injection</td>
<td>2018</td>
<td>\cite{26}</td>
<td></td>
</tr>
<tr>
<td>Goats</td>
<td><em>BLG</em> KO</td>
<td><em>BLG</em> abolished</td>
<td>CRISPR/Cas9 and zygote injection</td>
<td>2017</td>
<td>\cite{25}</td>
<td></td>
</tr>
<tr>
<td>Goats</td>
<td><em>BLG</em>-KO and <em>hLF</em>-KI</td>
<td>BLG-free and high- <em>hLF</em> milk</td>
<td>TALEN/SCNT</td>
<td>2015</td>
<td>\cite{35}</td>
<td></td>
</tr>
<tr>
<td>Meat</td>
<td>Cattle</td>
<td><em>MSTN</em> biallelic mutations</td>
<td>Double-muscled phenotype</td>
<td>ZFNs/SCNT</td>
<td>2014</td>
<td>\cite{4}</td>
</tr>
<tr>
<td>Goats</td>
<td><em>MSTN</em> biallelic KO</td>
<td>Disruption of <em>MSTN</em> expression</td>
<td>CRISPR/Cas9 and SCNT</td>
<td>2014</td>
<td>\cite{24}</td>
<td></td>
</tr>
<tr>
<td>Goats</td>
<td><em>MSTN</em>-KO and fat-1 KI</td>
<td>Muscle fibers stronger</td>
<td>CRISPR/Cas9 and SCNT</td>
<td>2018</td>
<td>\cite{52}</td>
<td></td>
</tr>
<tr>
<td>Sheep</td>
<td><em>MSTN</em>-KO</td>
<td>Muscle fibers stronger</td>
<td>CRISPR/Cas9 and SCNT</td>
<td>2018</td>
<td>\cite{53}</td>
<td></td>
</tr>
</tbody>
</table>
about by extended calving intervals, extra work for veterinarian and herdsman, increased culling rate and mortality risk, and reduced milk yield and quality[58]. The disease, caused by *Staphylococcus aureus*, has a low cure rate with antibiotics (often less than 25%) and there is currently no effective vaccine[59]. However, lysostaphin enables specific targeting of *S. aureus* through cleaving the pentaglycine bridge of the peptidoglycan of the bacterium[59]. Researchers had previously reported the generation of eight transgenic heifers, five of which survived into adulthood and secreted lysostaphin[60]. The highest expressing cows were not infected by intramammary infusion of *S. aureus*. However, the lysostaphin gene was randomly integrated into the bovine genome. Researchers reported ZFN nickases-mediated gene-edited cows with addition of the lysostaphin gene at the endogenous β-casein locus. The milk contained lysostaphin that was analyzed to confirm its ability to kill *S. aureus*[27]. Although comparison of treatment of *S. aureus* bovine mastitis with lysostaphin and a commonly used antibiotic revealed similar cure rates[61], it is feasible to protect gene-edited cows from mastitis by high expression of lysostaphin in milk.

Prion diseases are transmissible and fatal neurodegenerative disease[62] (such as bovine spongiform encephalopathy in cattle[63], and scrapie in sheep and goats[64]) that occur because of the conversion of the physiological cellular prion protein (PrPc) into the pathogenic β-sheets enriched isoform designated PrPSc and the propagation of PrPSc by recruiting PrPc[65]. It has been reported that mice devoid of PrPc were resistant to scrapie[66]. To test the feasibility of this strategy in cattle, 12 PrPc-deficient cattle were produced by HR-mediated KO method and SCNT technology[49]. Two PrPc-deficient cattle were subjected to histopathological analyses and showed no obvious abnormalities or significant lesions in any tissues. Protein misfolding cyclic amplification assay demonstrated the ability of the brain homogenates from one gene-edited individual to resist prion propagation. At over 20 months of age, the other eight cattle were still healthy. In addition, four PrPc-lambs were born, unfortunately three died soon after birth and one survived for 12 days[60]. The production of PrPc-deficient ruminants has assumed tremendous importance for disease resistance breeding and the production of prion-free biometrical materials.

Leukotoxin secreted by *Mannheimia hemolytica* causes acute inflammation and lung tissue damage[67]. Researchers using CRISPR/Cas9 introduced Q(-5)G substitution in both alleles of CD18 in bovine fetal fibroblasts, which impacted on leukotoxin binding to signal peptide of ruminant CD18 and abolished cytolysis of leukocytes in gene-edited cattle[511]. Also, researchers reported on the site-specific KI of the Cas9 nickase-mediated NRAMP1 gene and TALE nickase-mediated SP110 nuclear body protein gene to produce tuberculosis-resistant cattle[47]. These disease-resistant gene-edited cattle enabled the significant decrease in economic losses, maintaining animal health and reducing the use of antibiotics and prophylactics[60].

### 4.3 Animal welfare

Physical dehorning is a common method used to protect dairy cattle and humans from harm but results in reduced animal welfare and high cost. The development of SCNT and EENs mediated genome editing technology enabled the generation of hornless dairy cattle. In 2016, researchers achieved introgression of a putative POLLED allele into the genome of bovine embryo fibroblasts through TALEN-mediated integration and produced two healthy, homozygous polled dairy cattle by SCNT[31]. Furthermore, researchers simulated introgression of the POLLED allele into cattle populations via conventional breeding or gene editing over the course of 20 years in the USA[68]. In comparison to crossbreeding, gene editing enabled fast or faster decrease of frequency of the horned allele to < 0.1, a significant improvement of lifetime net merit and less inbreeding, which provided scientists with a powerful tool to provide polled genetics to the dairy industry. These reports illustrate the potential of genome editing in the production of polled cattle. However, a recent report by US Food and Drug Administration (FDA) scientists revealed the undesired heterozygous integration of bacterial DNA and a second copy of the repair template sequence in the genome of one of the edited polled cattle, although it had been claimed to be 100% bovine[69,70]. To reduce the risk of unintended consequences from gene-edited animals, it is thus important to improve the specificity of EENs[71,72] and select appropriate methods, such as long-range PCR, quantitative PCR, Southern blot, long-read sequencing[70] and unbiased genome-wide off-target detection[73,74], to measure plasmid integration events and off-target mutagenesis.

### 4.4 Milk and meat

Currently, we face the situation where the demand for animal products outstrips supply. Through EENs, gene-edited ruminants could improve the quality of milk and meat that are indispensable as a source of food for humans. BLG, considered to be one of the most important allergens in milk, cannot be abolished by heat treatment or fermentation[75,76]. In addition, BLG can cause allergic reactions even after gastrointestinal digestion due to its resistance to acid and protease hydrolysis[23]. The first report of gene-edited ruminants produced using EENs involved a cow containing biallelic mutations of the *BLG* gene which survived for over six months[22]. In 2018, researchers generated a ZFN-mediated marker-gene-free cloned cow, in which the *BLG* biallelic mutations consisted of 17-bp and 16-bp indels that could lead to frameshift mutations, resulting in BLG-free milk that caused lower
allergic reactions in Balb/c mice\textsuperscript{[23]}. Furthermore, researchers reported that BLG protein was abolished in cattle edited by zygote-mediated editing using TALEN\textsuperscript{[26]} and goats edited by co-injection of Cas9 mRNA and sgRNAs into one-cell-stage embryos\textsuperscript{[25]}. In 2015, researchers utilized TALEN to target the BLG locus to achieve KI of the hLF gene in fetal fibroblasts from goats\textsuperscript{[35]}. Analysis of the milk from the gene-edited goats revealed large-scale hLF expression and decreased expression of BLG.

Double-muscled animals have an increase in muscle mass of about 20\% because of a mutation of myostatin (MSTN)\textsuperscript{[77]}, which provides a direction for improving meat quality and economic benefit in ruminants. Researchers generated marker-gene-free MSTN biallelic mutation cattle via ZFNs combined with SCNT technology, which exhibited the double-muscled phenotype with about half the level of MSTN protein with an intact C-terminal domain estimated by ELISA analysis\textsuperscript{[4]}.

5 Discussion

SCNT technology enables the high efficiency of gene editing and the consistent expression of the target gene in gene-edited offsprings\textsuperscript{[78]} and permits full use of preferred germplasm for ruminant breeding\textsuperscript{[79,80]}. However, it still needs to be improved to overcome some problems, probably because of inbreeding between clones and incomplete epigenetic reprogramming of the somatic donor nuclei\textsuperscript{[23]}. For gene-edited cattle, compared to microinjection, SCNT requires more hours of practice and shows higher rates of midwifery and lower production efficiency. However, it is difficult to produce gene-edited calves via microinjection for monotocous cattle, because of chimerism and low numbers of zygotes. Thus, SCNT is currently the prevailing method to produce gene-edited cattle.

Random integration-mediated transgenic technology faces great safety and ethical problems because of its ability to generate uncontrolled gene transfer at unwanted sites and unregulated number of copies, which makes it easy to generate mutations of endogenous genes, to silence exogenous genes, and to integrate bacterial constituents into the target genome. HR-mediated gene editing technology could achieve exogenous DNA insertion to the target sites, which means transgenic technology is moving from an era of random integration into an era of site-specific integration. However, this is a huge challenge for large-scale production of gene-edited ruminants because of their low fertility. Hopefully, full development of the EENs mediated site-specific integration will significantly increase safety and efficiency of genome editing, which will greatly facilitate the production of gene-edited ruminants to serve human needs. Meanwhile, some optimizations have been developed for HDR-based KI technology including the quest for a double cut HDR donor\textsuperscript{[81]} and the optimal frequency at which HDR takes place\textsuperscript{[82]}.

Relying on the bacterial expression system for the production of certain proteins is most often impractical because of misfolding, aggregation of insoluble inclusion bodies and lack of posttranslational modifications\textsuperscript{[83]}. The use of insect cells, yeast and plants for therapeutic proteins also shows differences in glycosylation patterns, so are sometimes not suitable for therapeutic use in humans\textsuperscript{[84]}. Mammalian expression systems provide another option for the production of biomedical materials, for example with most of the current mAbs\textsuperscript{[85]}. However, low yield and high cost limit wider adoption\textsuperscript{[86]}. Compared to other species, cattle, sheep and goats have higher milk yield with existing dairy infrastructure\textsuperscript{[78]}. Therefore, gene-edited ruminants might be useful for efficiently and cheaply producing large and complex bioactive proteins, small unstable peptides, and proteins that require extensive posttranslational modification and are multimeric in nature\textsuperscript{[85]}. In addition to gene-edited animal models as bioreactors, both disease models of cystic fibrosis\textsuperscript{[87]} and human hypophosphatemia\textsuperscript{[88]} have been accomplished using CRISPR/Cas9 in sheep, which revealed characteristics consistent with disease symptoms observed in human patients and provided a novel large animal platform for human diseases.

Transgenic Atlantic salmon\textsuperscript{[14]} were developed by AquaBounty Technologies (Maynard, MA, USA) in 1992\textsuperscript{[89]}. Although transgenic salmon represent a significant milestone, they were only recently approved in Canada, so will finally come onto the market as the first edible transgenic animal. Interestingly, three pharmaceutical proteins produced by transgenic animals (ATryn produced in goats\textsuperscript{[90]}, Ruconest in rabbits\textsuperscript{[91]} and Kanuma in chicken eggs\textsuperscript{[92]} ) have been approved by the European Medicines Agency (EMA) and the FDA for medical use. Marketing transgenic ruminant products is more likely to be approved than the animals themselves, which can be limited by a series of issues, such as appropriate regulation, potential impact on the industry sector and social acceptability\textsuperscript{[93]}. The social acceptance of such gene-edited ruminant would depend on the major premise that factors such as the safety, animal welfare and ethical concerns, were taken into account\textsuperscript{[94]}. Public consideration of ruminant breeding by genome editing is largely based on their views on ethics, including moral standing, handling of animals, animal dignity and a critical analysis of the underlying relationship\textsuperscript{[95]}. It is possible that public consideration will be an essential reference and direction to the research and government regulation of gene-edited ruminant breeding. Gene-edited ruminant breeding mediated by site-specific EENs is safer and more controlled with potentially better market acceptance, in contrast to the transgenic breeding. However, the recent
controversy over edited polled cattle\cite{68} suggests that many people will be unlikely to accept the use of foreign plasmid in the process of genome editing. Thus, NHEJ-mediated genome editing may be considered for the production of ruminants because of the similarity to naturally occurring mutations\cite{94}.

### 6 Conclusions

Although the production of gene-edited ruminants has made great progress toward benefiting humans, the development of genome editing technology in ruminants has been much slower than in model organisms. However, the enormous scope for developing gene-edited ruminants holds immense promise for the future.

#### Acknowledgements

This work was supported by the National Major Project for Production of Transgenic Breeding (2016ZX08007003). We thank the FASE team for editing the English draft of this manuscript.

#### Compliance with ethics guidelines

Mengke Yuan, Yuanpeng Gao, Jing Han, Teng Wu, Jingcheng Zhang, Yongke Wei, and Yong Zhang declare that they have no conflicts of interest or financial conflicts to disclose.

This article is a review and does not contain any studies with human or animal subjects performed by any of the authors.

### References

8. Capecci M R. How close are we to implementing gene targeting in animals other than the mouse? *Proceedings of the National Academy of Sciences of the United States of America*, 2000, 97(3): 956–957
55. Houdebine L M. Production of pharmaceutical proteins by transgenic animals. Comparative Immunology, Microbiology and Infectious Diseases, 2009, 32(2): 107–121
2000, 21(8): 364–370