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Popular artícle

applications of CRÍSPR/Cas9 Genome Editing Technology in Veterinary Medicine

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abstract

Genome edíting technology is a technique for targeted genetic modifications, enabling the knockout and knock in of specific DNa fragments in the selected genome. This technology has been widely used in various types of biomedical, clinical and agricultural research. Three gene editing techniques including Zinc-finger nucleases (ZFNs), Transcription activator-like effector nucleases (Talens), and Crispr/Cas9 are commonly used in biomedical and life science research, with Crispr/Cas9 now being the most widely used. In this review, we discuss the applications of Crispr/Cas9 genome editing in the fields of veterinary medicine and animal husbandry.

Íntroduction

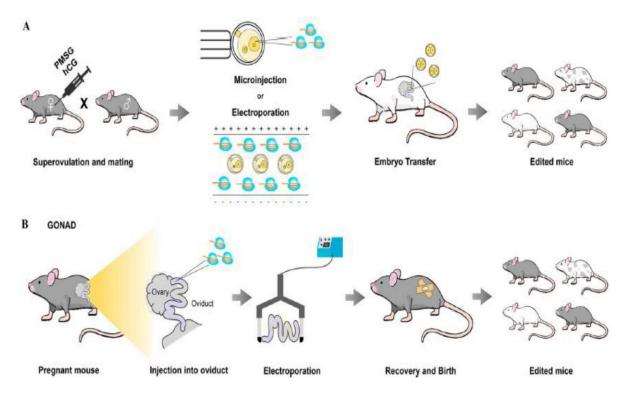
RÍSPR ís an acronym for "Clustered Regularly Ínterspaced Short Palíndromíc Repeats." CRÍSPR genome engíneeríng technology allows researchers to easíly edít the DNa of any genome. Naturally, the CRÍSPR/Cas system plays an important role in microbíal immunity. Ít acts as self-defence system in a sequence specific manner against exogenous virus or plasmíd in bactería by cleaving their DNa or RNa. When a virus or bactería infects a microbíal cell, the microbe employs a special CRÍSPR-associated nuclease (Cas9) to chop off a piece of the foreign DNa. The short RNa fragment known as a guide RNa (gRNa) directs the nuclease to its target sequence. The chopped off DNa fragment maybe then stored between the palindromíc CRÍSPR sequences to retain a genetic memory for disabling future infections from the same viral strain (Hílle and Charpentier, 2016).

applications of CRÍSPR/Cas9 genome editing technology in veterinary medicine and animal husbandry

The use of CRÍSPR/Cas9 technology in the field of veterinary research is greatly revolutionizing the ability to manipulate the animal genome to create appropriate disease models and disease resistant animals and also improves the quality meat production and animal welfare.

Transgenic animal models development

Developing appropriate animal models is necessary to understand the pathobiology and molecular mechanisms of human and animal diseases and it also plays an important role in drug development and organ transplantation. Many animal disease models have been generated for basic and clinical research by combining reproductive technologies like micro injection with genome editing. In addition, these animal models play an important role in the field of preclinical gene therapy and stem cell therapy research for rare genetic diseases.



Fígure 1. a: Transgeníc míce productíon usíng CRÍSPR system. a. CRÍSPR delívery to embryos usíng mícroínjectíon or electroporatíon. Edíted 2-cell-stage embryos are transplanted into a surrogate mother and the genome edíted offspring are obtained. **B:** Direct injection of the Cas9/gRNa complex into oviduct of pregnant mouse, followed by an electrical impulse produce genome edited pups (adopted from Lee et al. 2020).

Table 1. List of animal models developed by using CRÍSPR/Cas9 technique (Lee et al. 2020)

Species	Dísease model	Targeted gene	Techníque
	Tyrosínemía	Fah	Mícroínjectíon
Míce	Hemophílía a	F8	íPSC correctíon & ÍV injection
	Hemophílía B	F9	Íntravenous ínjectíon
	Duchenne Muscular Dystrophy	Dystropín	Íntramuscular ínjectíon
Rat	Retinal dystrophy	Rho	Subretinal injection
	Paríkínson's dísease	TH	Íntracraníal ínjectíon
Píg	Líver faílure, traumatíc shock	alb	Mícroínjectíon
	Huntíngton's dísease	Huntíngtín	Somatíc cell nuclear transfer
Dog	Muscular Dystrophy	Dystropín	Íntramuscular ínjectíon

Cancer biology

Since the cancer genome is highly complex, with hundreds of point mutations, translocations, and chromosomal aberrations per tumour, suitable animal models are needed to understand the effects and mechanisms of these alterations. Traditional methods to develop mouse models for oncological studies are time-consuming and laborious. The recent development of the CRÍSPR-Cas9 system is improving the generation of mouse models to study cancer biology.

Table 2. List of mice cancer models developed by using CRÍSPR/Cas9 (Mou et al. 2015)

Cancer model	Target tíssue	Delívery	Genes targeted
Rett syndrome	Embryo	One-cell embryo injection	Mecp2 Cre-LoxP
Colon cancer	ES cell	Íntra perítoneal	p53, apc, Pten
Hepatocellular cancer	Líver	Hydrodynamíc ínjectíon	Pten, p53, β-catenín

acute Myeloid	Fetal-líver	Íntravenous ínjectíon of	MII3
Leukemía	HSCs ex vívo	Cas9-edíted HSCs	
lung cancer	Lung	Íntranasal/íntra-tracheal	p53 and Lkb1, Kras
Dox-índucíble Burkít	Fetal-líver	Íntravenous ínjectíon of	Mcl-1, p53
lymphoma model	HSCs ex vívo	Cas9-edíted HSCs	

Swine production and research

Pígs are important domestic animals reared for food and pharmaceutical applications; they also served as ideal animal models for various human diseases such as diabetes, obesity, atherosclerosis and other cardiovascular diseases. Pork is an important meat source in the western countries. Worldwide, pig production accounted for 42% of total livestock production in 2018, and this percentage is expected to increase in the coming years. However, the existing breeding methods are not enough to meet the developing needs of pig production. The use of CRÍSPR-Cas9 technique has greatly promoted the advancement of pig production and research (Lín et al. 2019).

Following are the applications of CRÍSPR/Cas9 genome editing technique in the field of pig breeding and research:

- 1. Ín the development of rapíd víral vaccínes agaínst píg pathogens such as pseudorabíes vírus, porcíne reproductíve and respíratory syndrome vírus, classícal swíne fever vírus and afrícan swíne fever vírus.
- 2. Ín fast and reliable breeding and reproduction of disease resistant pigs. Ín 2017, Whitworth et al. used CRÍSPR/Cas9 technique to generate CD163-knockout pigs to protect pig from porcine reproductive and respiratory syndrome virus (PRRSV).
- 3. Ín the fíeld of transplant ímmunology studíes whích use pígs as an anímal model. Sato et al. ín 2013 and Petersen et al. ín 2016 created píglets with biallelíc knockouts of GGTa1 gene by usíng the CRÍSPR/Cas9 system. Ít was proved to be a best model to study xenotransplantation.
- 4. Ín the development of swíne dísease models used ín the translatíonal medícal research. Swíne models of human type Í and ÍÍÍ *von* Wíllebrand dísease, Huntíngton's dísease; ínsulín-defícient pígs for díabetes research, RUNX3-associated stomach cancer have been developed so far by usíng CRÍSPR/Cas9 technology.

5. CRÍSPR/Cas9 technology also improved the quality of pork. amount of fat and lean meat contents are important factors determines the palatability of pork. Myostatin (MSTN) knockout cloned pigs without selectable marker gene (SMG) developed by combined use of CRÍSPR/Cas9 and Cre/LoxP showed more pronounced skeletal muscles and decreased back fat thickness.

Farm animal production and research

Genome edíting in the farm animals (bovine) is majorly focused on the production of disease resistant animals (e.g., tuberculosis, brucellosis), improved generation of meat and dairy products, animal sexing, introduction of desirable phenotypes (e.g., stress tolerance, disease resistance) and animal welfare (e.g., polled or hornless).

Following are the applications of CRÍSPR/Cas9 genome editing technique in the field of farm animal breeding and research:

1. Production of disease resistant animals

Tuberculosís-resístant genetícally modífied cattle (NRaMP1 knock-ín), bovíne spongíform encephalopathy, and chroníc wasting dísease resístant cattle (PRNP knock-out), Jhone's dísease resístant cattle (ÍL10Ra knock-out), and brucellosís resístant cow (vírB10 or Rpola transductíon) were generated by CRÍSPR/Cas9 medíated genomíc edíting of bovíne genome (Síngh and alí, 2021).

2. Ímprovíng anímal Welfare

a horned phenotype of bovíne (plethora) íncreases the rísk of ínjury or damage to the anímal and handler. a polled (hornless) phenotype ís preferred ín thís case. Usually, polled phenotypes are essentíally used ín angus meat breeds. Ín dífferent meat breeds, the polled Celtíc (Pc) variation, within the polled locus, índuces a polled phenotype. Schuster et al. ín 2020 produced polled HF bulls by incorporating the Pc variety into its genome by using CRÍSPR/Cas12a framework which eliminated the need for dehorning.

3. Ímprovíng semen sexíng

Semen sexing to find out the sex of developing embryos before foetus transition in animals was improved by knock-in eGFP (green fluorescent protein) gene within the Y-chromosome of bovine fetal fibroblast (BFF) cell lines with the assistance of CRÍSPR/Cas9 (Zhao et al. 2020).

Conclusion

Over the past few years, CRÍSPR/Ca9 genome-edíting technology improved the development of genetically engineered animals which could be served as animal models for various human diseases in translational research. The use of the CRÍSPR/Ca9 technique in veterinary and animal husbandry research showed promising results in the production of disease resistant animals, quality meat and also improved the animal welfare. This novel technology will continue to revolutionize veterinary medicine. Precision animal models will pave the way for precision drug discovery.

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