

# Zscan26 Cas9-KO Strategy

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# **Project Overview**



**Project Name** 

Zscan26

**Project type** 

Cas9-KO

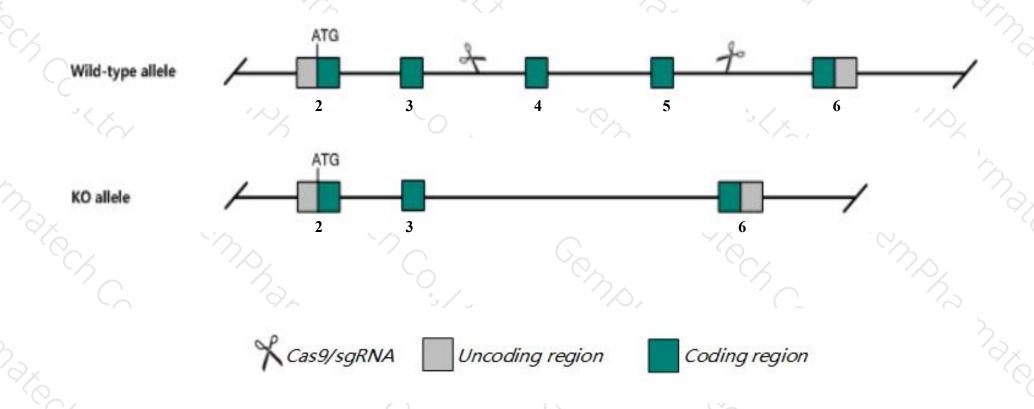
Strain background

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Zscan26 gene. The schematic diagram is as follows:



### **Technical routes**



- The Zscan26 gene has 2 transcripts. According to the structure of Zscan26 gene, exon4-exon5 of Zscan26-202(ENSMUST00000110485.2) transcript is recommended as the knockout region. The region contains 427bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zscan26* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6JGpt mice.Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

### **Notice**



- > The Zscan26 gene is located on the Chr13. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

## Gene information (NCBI)



#### Zscan26 zinc finger and SCAN domain containing 26 [Mus musculus (house mouse)]

Gene ID: 432731, updated on 13-Mar-2020

#### Summary

△ ?

Official Symbol Zscan26 provided by MGI

Official Full Name zinc finger and SCAN domain containing 26 provided by MGI

Primary source MGI:MGI:3531417

See related Ensembl:ENSMUSG00000022228

Gene type protein coding
RefSeq status VALIDATED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as BC068174, Zfp187, Znf187

Expression Ubiquitous expression in bladder adult (RPKM 25.1), cerebellum adult (RPKM 17.0) and 25 other tissuesSee more

Orthologs <u>human all</u>

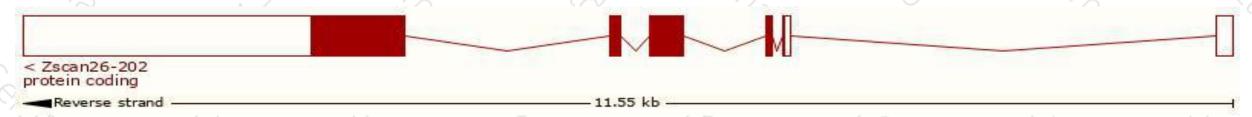
# Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

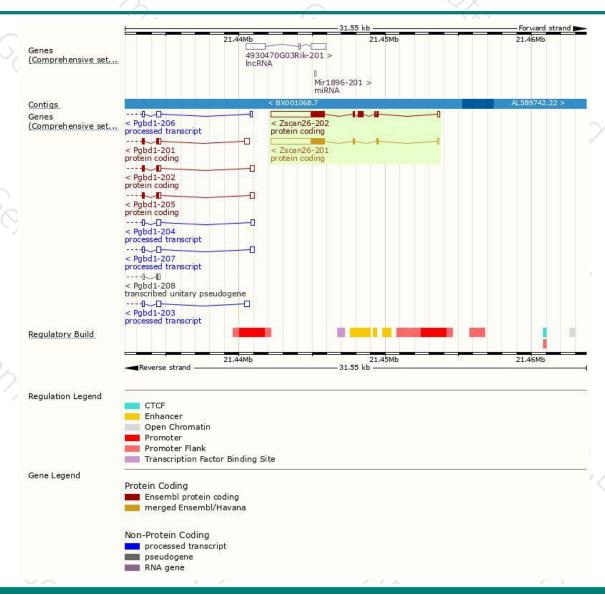
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zscan26-202	ENSMUST00000110485.2	4366	466aa	Protein coding	CCDS84009	Q5RJ54	TSL:5 GENCODE basic APPRIS P1
Zscan26-201	ENSMUST00000032820.14	3918	<u>340aa</u>	Protein coding	CCDS49211	F8WJ31	TSL:3 GENCODE basic

The strategy is based on the design of *Zscan26-202* transcript, the transcription is shown below:



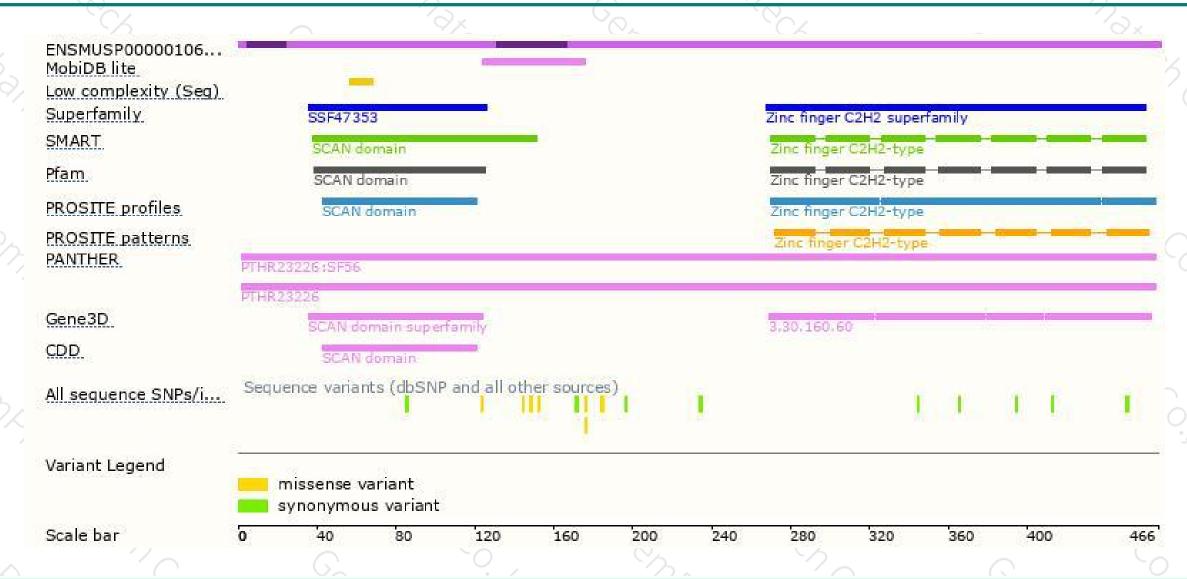
### Genomic location distribution





### Protein domain







If you have any questions, you are welcome to inquire.

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