

# Zfp189 Cas9-CKO Strategy

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# **Project Overview**



**Project Name** 

Zfp189

**Project type** 

Cas9-CKO

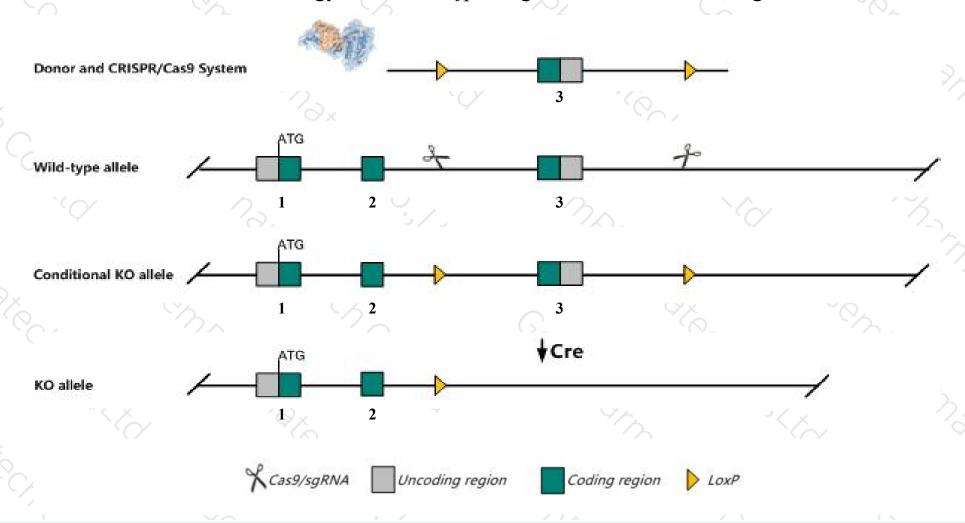
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Zfp189 gene. The schematic diagram is as follows:



### Technical routes



- The Zfp189 gene has 2 transcripts. According to the structure of Zfp189 gene, exon3 of Zfp189-202(ENSMUST00000107696.1) transcript is recommended as the knockout region. The region contains 1709bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zfp189* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA and Donor 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.
- > The flox mice was knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

### **Notice**



- > The Zfp189 gene is located on the Chr4. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

### Gene information (NCBI)



#### Zfp189 zinc finger protein 189 [Mus musculus (house mouse)]

Gene ID: 230162, updated on 13-Mar-2020

#### Summary

☆ ?

Official Symbol Zfp189 provided by MGI

Official Full Name zinc finger protein 189 provided by MGI

Primary source MGI:MGI:2444707

See related Ensembl: ENSMUSG00000039634

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 BB131811, C430015I23Rik

Expression Ubiquitous expression in CNS E11.5 (RPKM 2.7), CNS E14 (RPKM 2.5) and 25 other tissuesSee more

Orthologs <u>human all</u>

# Transcript information (Ensembl)



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

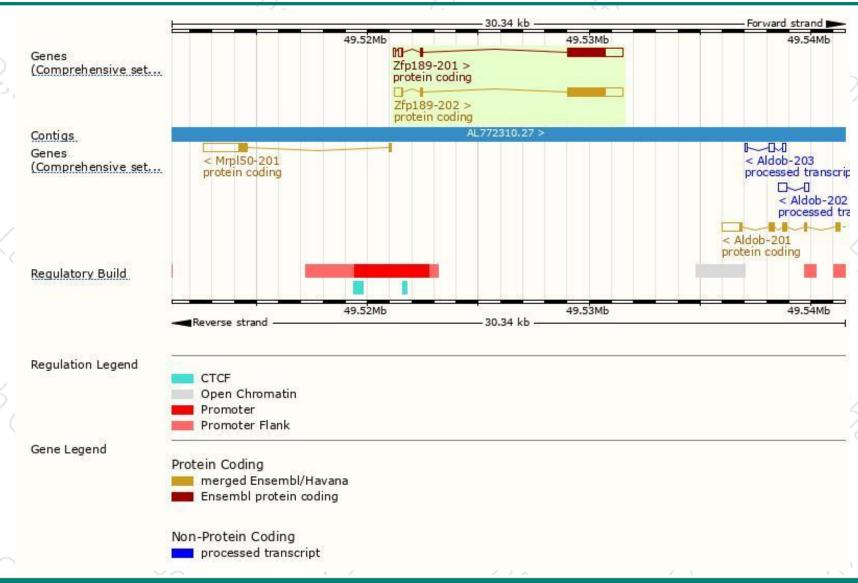
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp189-202	ENSMUST00000107696.1	2965	608aa	Protein coding	CCDS18175	Q8BKP2	TSL:1 GENCODE basic APPRIS P1
Zfp189-201	ENSMUST00000042964.12	2835	608aa	Protein coding	CCDS18175	Q8BKP2	TSL:2 GENCODE basic APPRIS P1

The strategy is based on the design of Zfp189-202 transcript, the transcription is shown below:



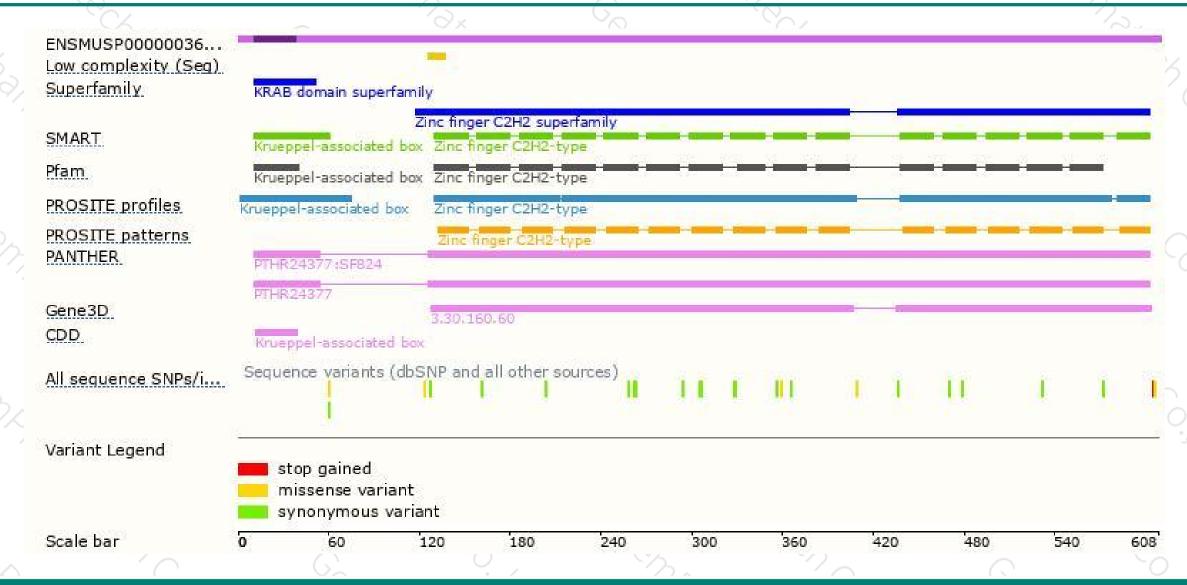
### Genomic location distribution





### Protein domain







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

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