

***Zfp7* Cas9-CKO Strategy**

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Project Overview

Project Name

Zfp7

Project type

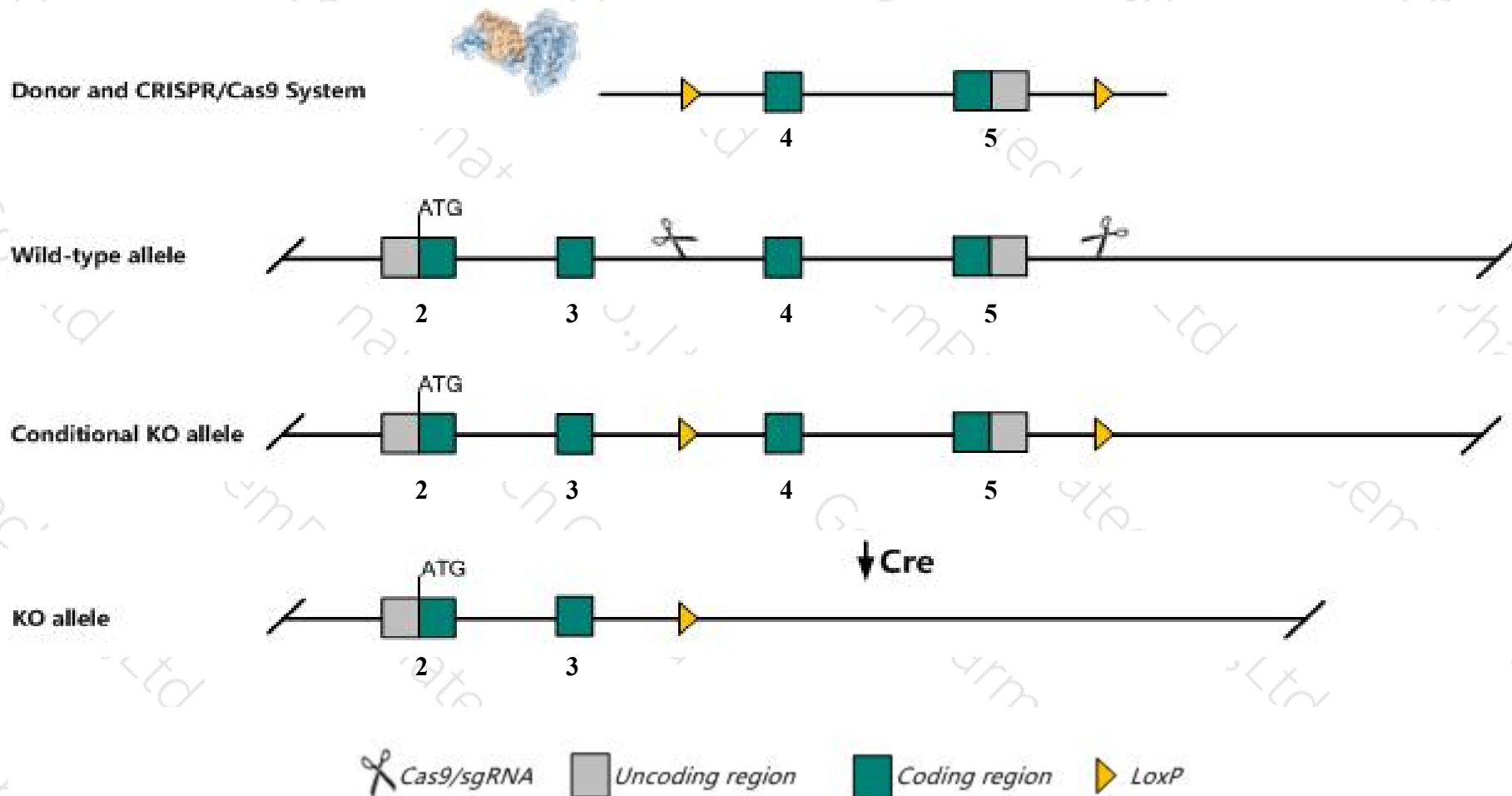
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zfp7* gene. The schematic diagram is as follows:



- The *Zfp7* gene has 6 transcripts. According to the structure of *Zfp7* gene, exon4-exon5 of *Zfp7*-205(ENSMUST00000230106.1) transcript is recommended as the knockout region. The region contains 1928bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zfp7* 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 *Zfp7* gene is located on the Chr15. 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)

Zfp7 zinc finger protein 7 [Mus musculus (house mouse)]

Gene ID: 223669, updated on 13-Mar-2020

Summary



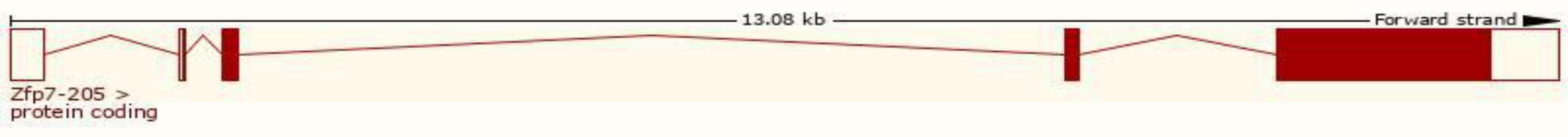
Official Symbol	Zfp7 provided by MGI
Official Full Name	zinc finger protein 7 provided by MGI
Primary source	MGI:MGI:99208
See related	Ensembl:ENSMUSG00000033669
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	KRAB20, KRAB7, Krox-2, Zfp-7, Zfp65, Zfp80, Zfp86-rs1, mszf73-2
Expression	Ubiquitous expression in CNS E11.5 (RPKM 4.7), bladder adult (RPKM 4.6) and 27 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

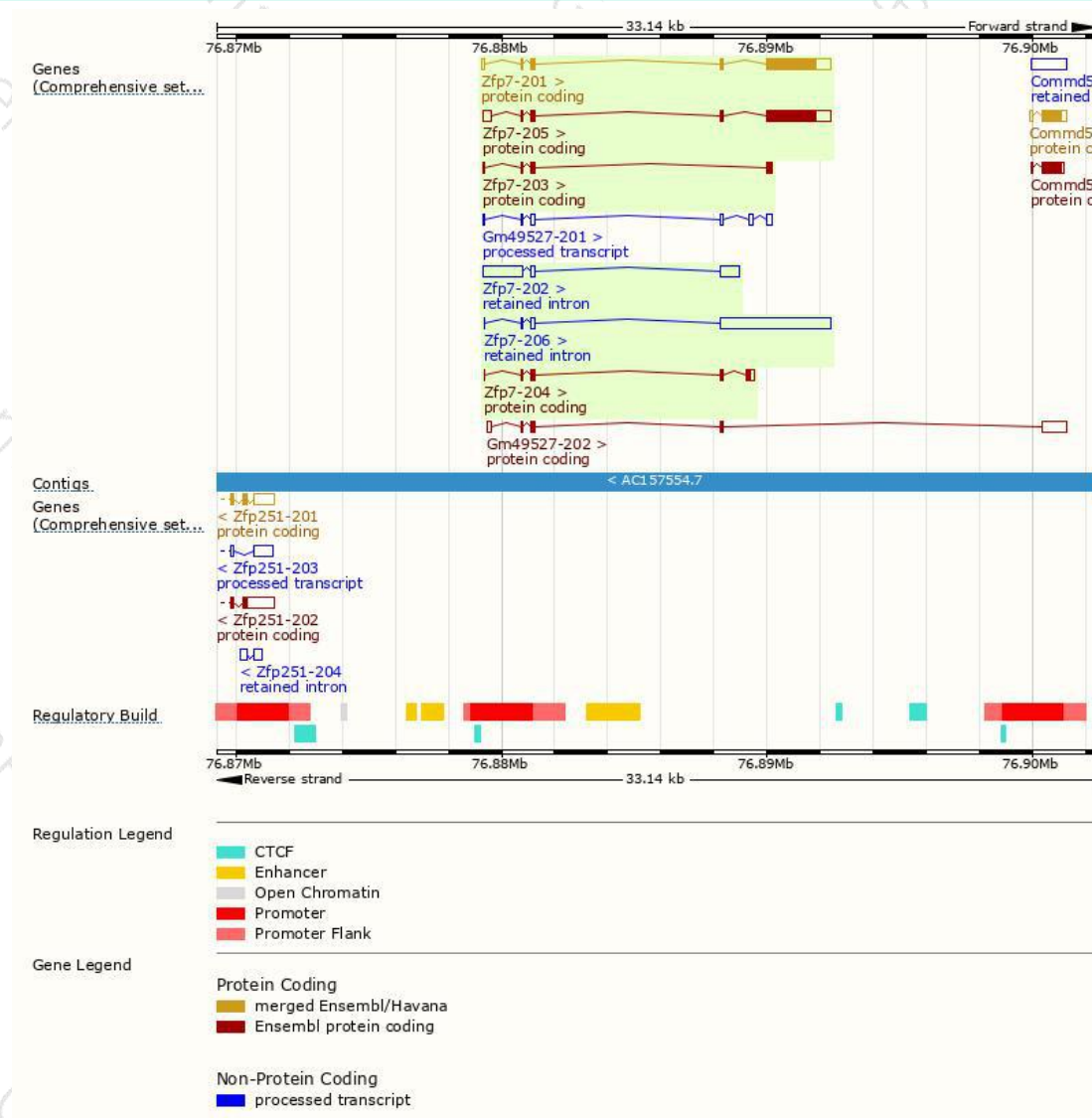
The gene has 6 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp7-205	ENSMUST00000230106.1	2960	685aa	Protein coding	CCDS37124	Q3TFZ4	GENCODE basic APPRIS P1
Zfp7-201	ENSMUST0000023179.6	2772	685aa	Protein coding	CCDS37124	Q3TFZ4	TSL:1 GENCODE basic APPRIS P1
Zfp7-204	ENSMUST00000229990.1	644	131aa	Protein coding	-	A0A2R8VHC1	GENCODE basic
Zfp7-203	ENSMUST00000229831.1	395	103aa	Protein coding	-	A0A2R8VI83	CDS 3' incomplete
Zfp7-206	ENSMUST00000230954.1	4356	No protein	Retained intron	-	-	
Zfp7-202	ENSMUST00000229729.1	2340	No protein	Retained intron	-	-	

The strategy is based on the design of *Zfp7-205* 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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