

Zfp469 Cas9-CKO Strategy

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Project Overview

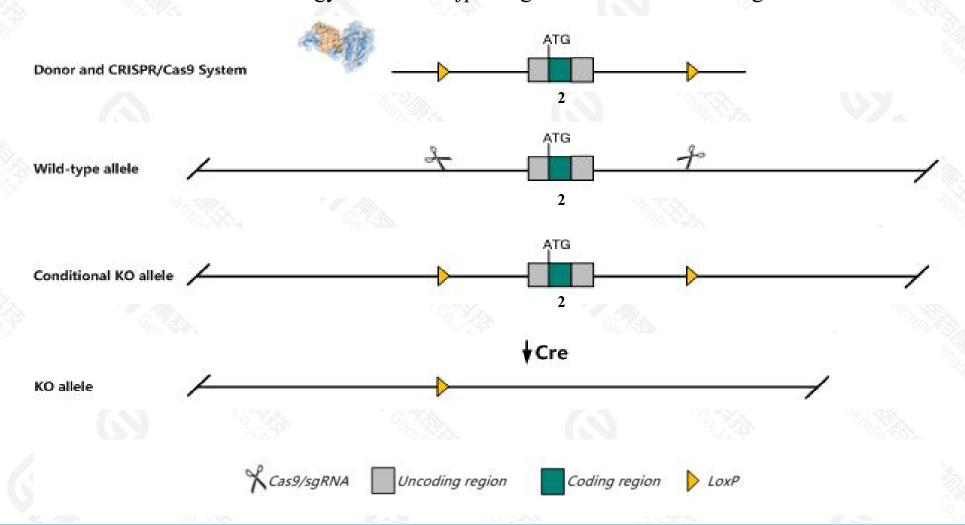


Project Name	Zfp469
Project type	Cas9-CKO
Strain background	C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Zfp469 gene. The schematic diagram is as follows:



Technical routes



- > The Zfp469 gene has 2 transcripts. According to the structure of Zfp469 gene, exon2 of Zfp469-202(ENSMUST00000187142.2) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zfp469* 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 KO region contains partial intron of the *Galnt2l* gene.Knockout the region may affect the function of *Galnt2l* gene.
- > The Zfp469 gene is located on the Chr8. 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)



Zfp469 zinc finger protein 469 [Mus musculus (house mouse)]

Gene ID: 195209, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Zfp469 provided by MGI

Official Full Name zinc finger protein 469 provided by MGI

Primary source MGI:MGI:2684868

See related Ensembl: ENSMUSG00000043903

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 Gm22

Expression Broad expression in ovary adult (RPKM 1.6), limb E14.5 (RPKM 1.6) and 19 other tissuesSee more

Orthologs <u>human</u> all

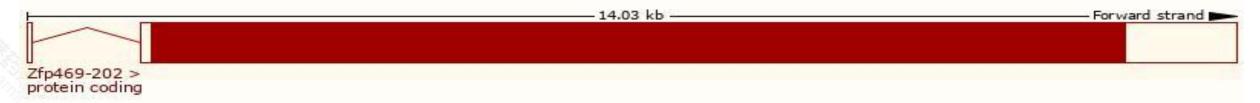
Transcript information (Ensembl)



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

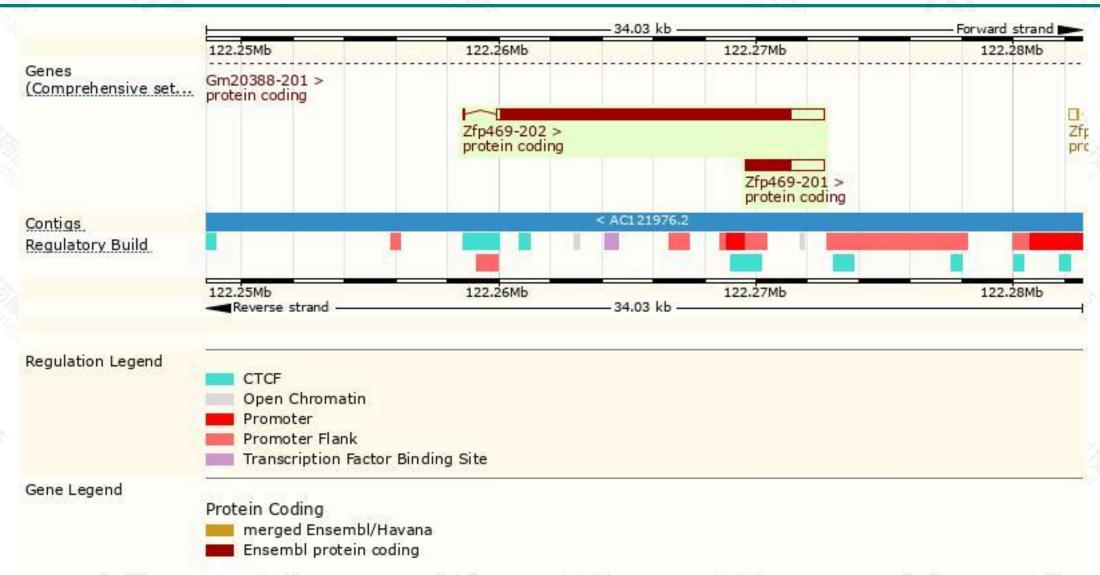
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp469-202	ENSMUST00000187142.2	12777	3765aa	Protein coding	-		TSL:NA GENCODE basic APPRIS P1
Zfp469-201	ENSMUST00000055537.2	3078	<u>597aa</u>	Protein coding	-		TSL:NA GENCODE basic

The strategy is based on the design of *Zfp469-202* 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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