

Fbxo28 Cas9-CKO Strategy

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Project Overview



Project Name

Fbxo28

Project type

Cas9-CKO

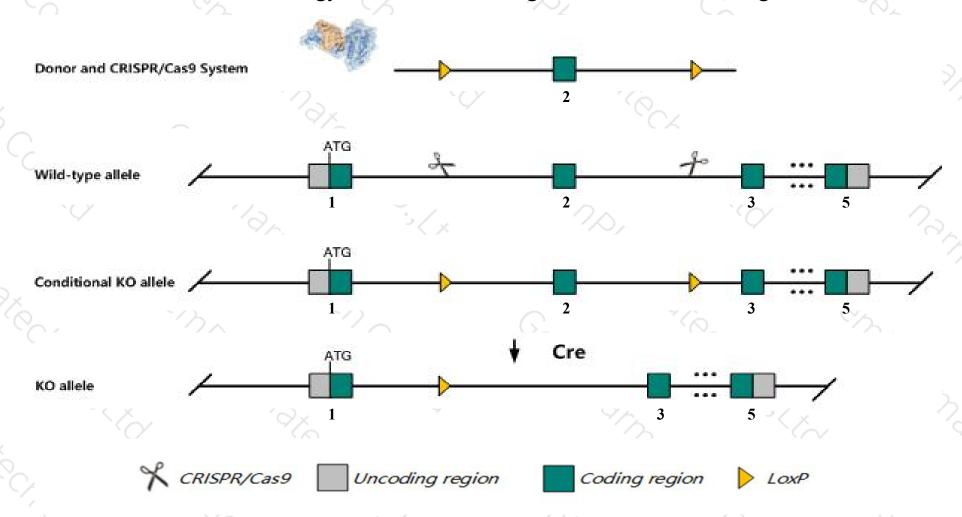
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Fbxo28* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Fbxo28* gene has 5 transcripts. According to the structure of *Fbxo28* gene, exon2 of *Fbxo28-201*(ENSMUST00000051431.9) transcript is recommended as the knockout region. The region contains 110bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fbxo28* gene. The brief process is as follows:CRISPR/Cas9 system 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 will be 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 *Fbxo28* gene is located on the Chr1. 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)



Fbxo28 F-box protein 28 [Mus musculus (house mouse)]

Gene ID: 67948, updated on 31-Jan-2019

Summary

△ ?

Official Symbol Fbxo28 provided by MGI

Official Full Name F-box protein 28 provided by MGI

Primary source MGI:MGI:1261890

See related Ensembl: ENSMUSG00000047539

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 4833428J17Rik, 5730505P19Rik, D1Ertd578e, Fbx28, mKIAA0483

Expression Ubiquitous expression in whole brain E14.5 (RPKM 7.0), CNS E14 (RPKM 6.8) and 28 other tissuesSee more

Orthologs <u>human</u> all

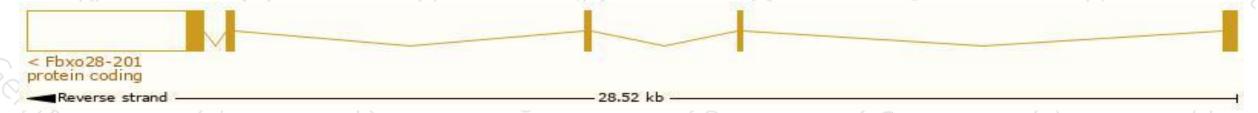
Transcript information (Ensembl)



The gene has 5 transcripts, all transcripts are shown below:

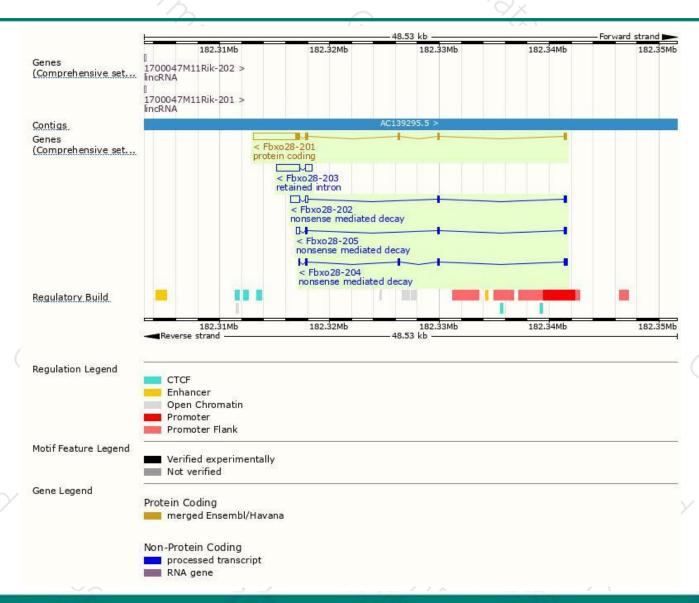
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fbxo28-201	ENSMUST00000051431.9	4925	368aa	Protein coding	CCDS15588	Q8BIG4	TSL:1 GENCODE basic APPRIS P1
Fbxo28-202	ENSMUST00000192544.5	1407	<u>126aa</u>	Nonsense mediated decay	-	A0A0A6YX52	TSL:5
Fbxo28-205	ENSMUST00000195061.5	781	<u>143aa</u>	Nonsense mediated decay	020	A0A0A6YX00	CDS 5' incomplete TSL:3
Fbxo28-204	ENSMUST00000194213.1	747	<u>195aa</u>	Nonsense mediated decay	798	A0A0A6YVQ6	TSL:3
Fbxo28-203	ENSMUST00000193700.1	2676	No protein	Retained intron	125)	-	TSL:1

The strategy is based on the design of Fbxo28-201 transcript, The transcription is shown below



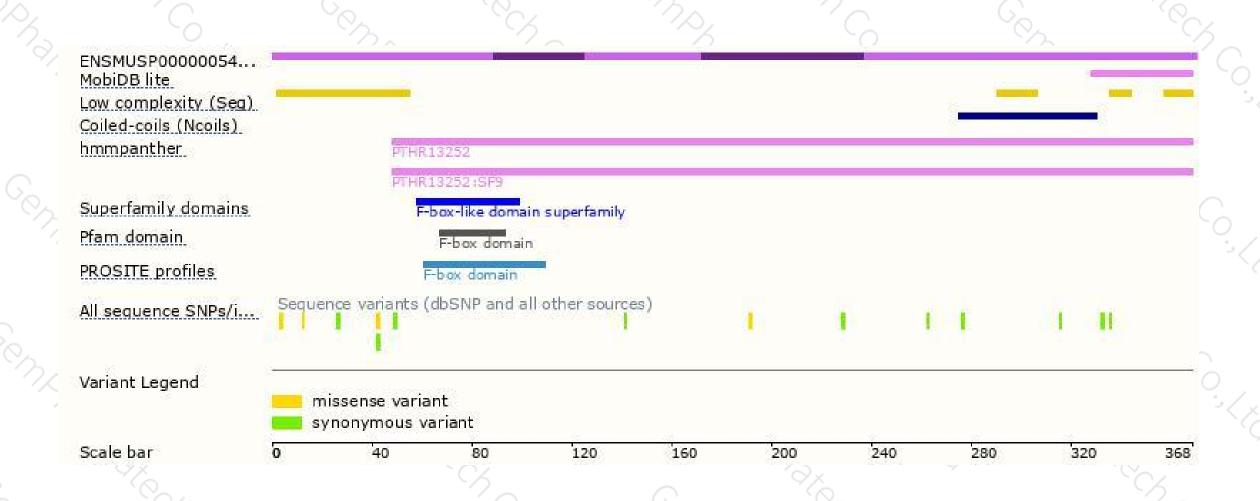
Genomic location distribution





Protein domain







If you have any questions, you are welcome to inquire. Tel: 400-9660890





