

Fbxo25 Cas9-KO Strategy

Designer: Yanhua Shen

Design Date: 2019-08-07

Project Overview



Project Name

Fbxo25

Project type

Cas9-KO

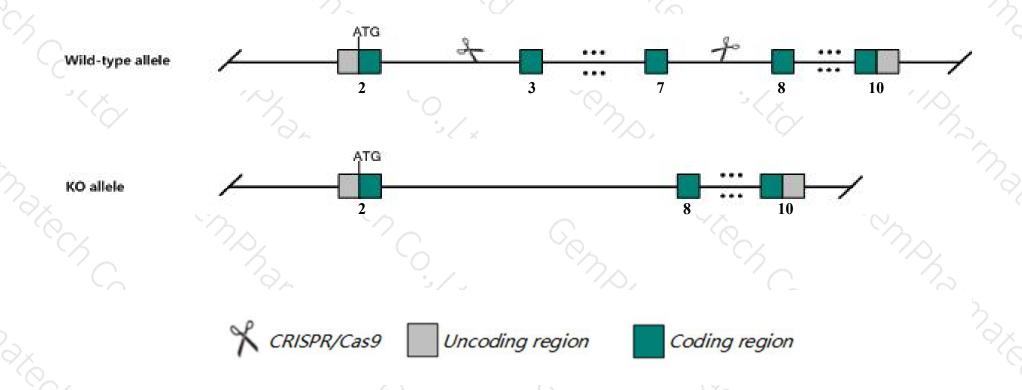
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Fbxo25* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Fbxo25* gene has 5 transcripts. According to the structure of *Fbxo25* gene, exon3-exon7 of *Fbxo25-201*(ENSMUST00000043520.4) transcript is recommended as the knockout region. The region contains 523bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify Fbxo25 gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ The *Fbxo25* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Fbxo25 F-box protein 25 [Mus musculus (house mouse)]

Gene ID: 66822, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Fbxo25 provided by MGI

Official Full Name F-box protein 25 provided by MGI

Primary source MGI:MGI:1914072

See related Ensembl: ENSMUSG00000038365

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 9130015106Rik, Al649137, Fbx25

Expression Ubiquitous expression in duodenum adult (RPKM 42.5), colon adult (RPKM 41.2) and 28 other tissuesSee more

Orthologs <u>human</u> all

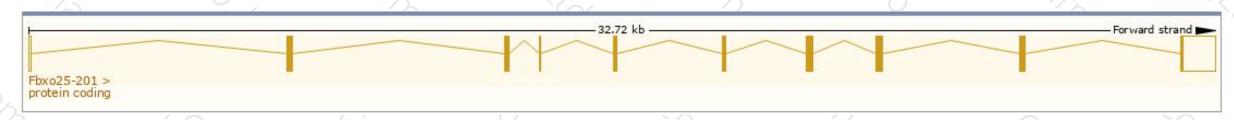
Transcript information (Ensembl)



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

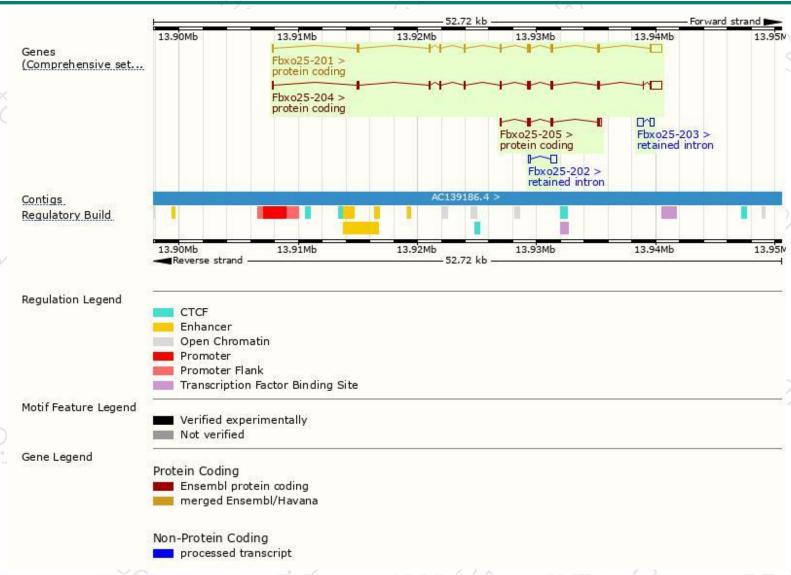
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fbxo25-204	ENSMUST00000209913.1	2035	<u>365aa</u>	Protein coding	CCDS85507	A0A1B0GRC8	TSL:1 GENCODE basic APPRIS ALT2
Fbxo25-201	ENSMUST00000043520.4	2012	<u>357aa</u>	Protein coding	CCDS22118	Q9D2Y6	TSL:1 GENCODE basic APPRIS P3
Fbxo25-205	ENSMUST00000210280.1	737	201aa	Protein coding	14	A0A1B0GSB9	CDS 5' incomplete TSL:3
Fbxo25-203	ENSMUST00000209310.1	781	No protein	Retained intron	i e	75 <u>2</u> 8	TSL:2
Fbxo25-202	ENSMUST00000209290.1	646	No protein	Retained intron		(2)	TSL:3

The strategy is based on the design of *Fbxo25-201* transcript, The transcription is shown below



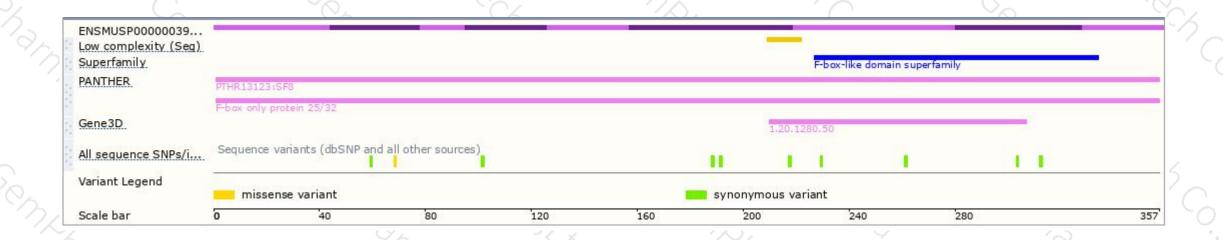
Genomic location distribution





Protein domain







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





