

Fbxo42 Cas9-KO Strategy

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Design Date: 2018-6-19

Project Overview



Project Name

Fbxo42

Project type

Cas9-KO

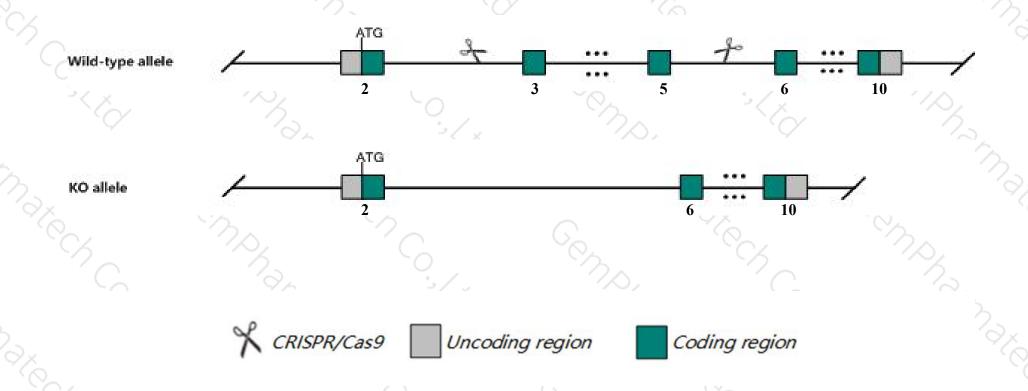
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Fbxo42* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Fbxo42* gene has 2 transcripts. According to the structure of *Fbxo42* gene, exon3-exon5 of *Fbxo42-201* (ENSMUST00000030757.9) transcript is recommended as the knockout region. The region contains 406bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Fbxo42* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



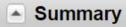
- ➤ The *Fbxo42* gene is located on the Chr4. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Fbxo42 F-box protein 42 [Mus musculus (house mouse)]

Gene ID: 213499, updated on 12-Aug-2019



Official Symbol Fbxo42 provided by MGI

Official Full Name F-box protein 42 provided by MGI

Primary source MGI:MGI:1924992

See related Ensembl:ENSMUSG00000028920

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 mKIAA1332; 6720460I06Rik

Expression Ubiquitous expression in thymus adult (RPKM 8.6), whole brain E14.5 (RPKM 8.0) and 28 other tissues <u>See more</u>

Orthologs human all

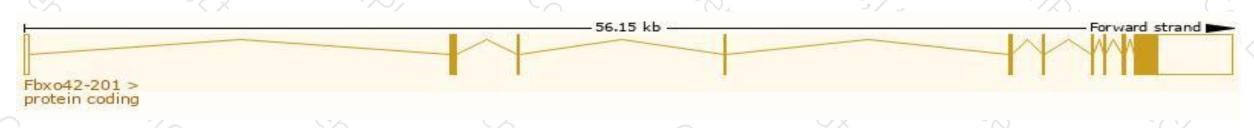
Transcript information (Ensembl)



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

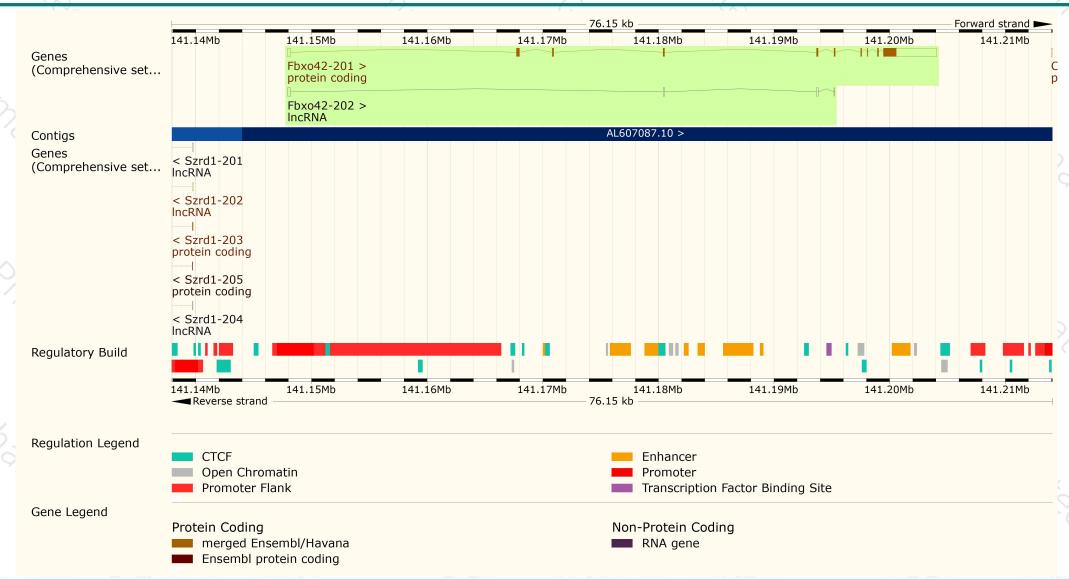
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fbxo42-201	ENSMUST00000030757.9	5934	<u>717aa</u>	Protein coding	CCDS18866	Q6PDJ6	TSL:1 GENCODE basic APPRIS P1
Fbxo42-202	ENSMUST00000146768.1	579	No protein	Processed transcript	N -		TSL:3

The strategy is based on the design of Fbxo42-201 transcript, The transcription is shown below



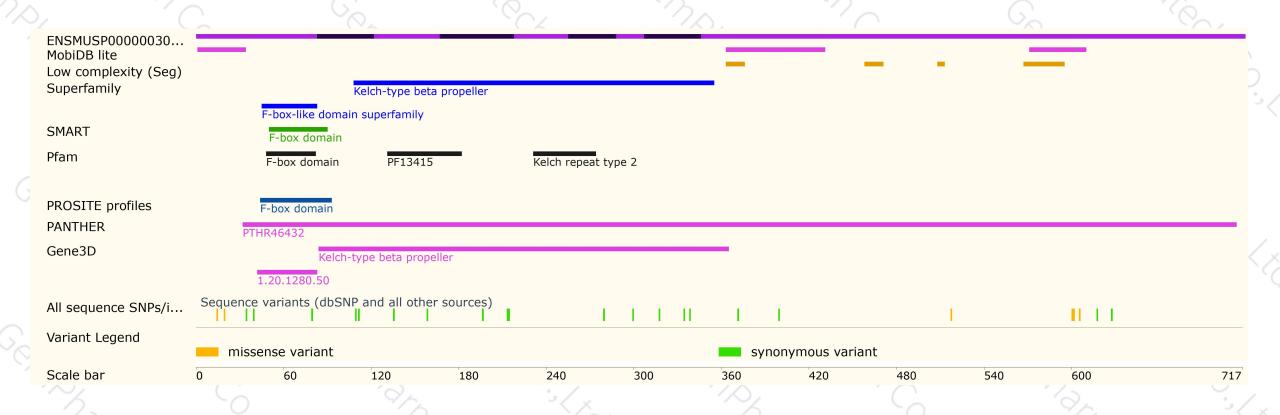
Genomic location distribution





Protein domain







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





