

Snrnp40 Cas9-KO Strategy

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Project Overview



Project Name

Snrnp40

Project type

Cas9-KO

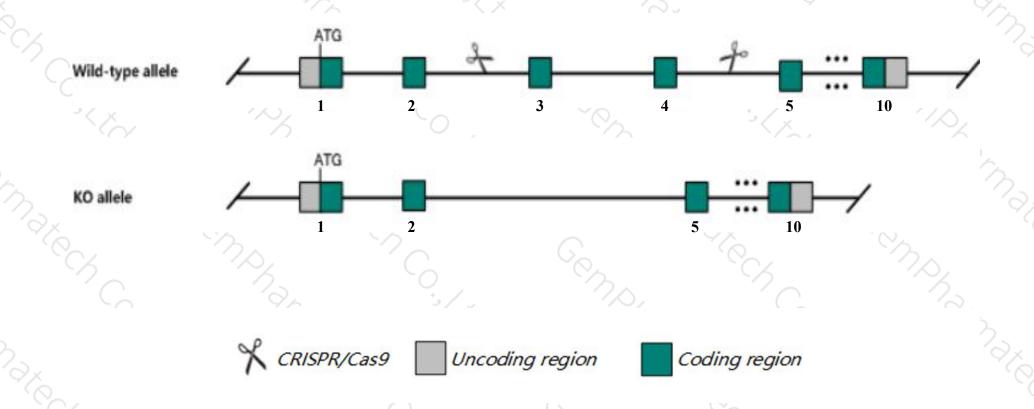
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Snrnp40* gene. The schematic diagram is as follows:



Technical routes



- > The Snrnp40 gene has 4 transcripts. According to the structure of Snrnp40 gene, exon3-exon4 of Snrnp40-201(ENSMUST00000105994.3) transcript is recommended as the knockout region. The region contains 260bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Snrnp40* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice



- > The Snrnp40 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)



Snrnp40 small nuclear ribonucleoprotein 40 (U5) [Mus musculus (house mouse)]

Gene ID: 66585, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Snrnp40 provided by MGI

Official Full Name small nuclear ribonucleoprotein 40 (U5) provided by MGI

Primary source MGI:MGI:1913835

See related Ensembl:ENSMUSG00000074088

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 0610009C03Rik, Prp8bp, Wdr57

Expression Ubiquitous expression in CNS E11.5 (RPKM 42.3), limb E14.5 (RPKM 31.2) and 28 other tissuesSee more

Orthologs <u>human all</u>

Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

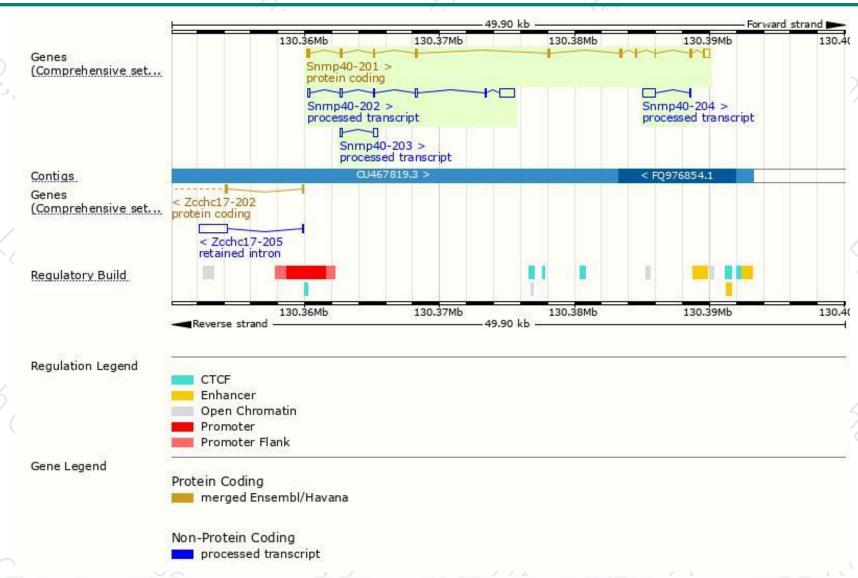
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Snrnp40-201	ENSMUST00000105994.3	1584	358aa	Protein coding	CCD538892	Q6PE01	TSL:1 GENCODE basic APPRIS P1
Snrnp40-202	ENSMUST00000180577.1	1752	No protein	Processed transcript	-	-8	TSL:1
Snrnp40-204	ENSMUST00000181560.1	1048	No protein	Processed transcript	-	20	TSL:5
Snrnp40-203	ENSMUST00000181161.1	489	No protein	Processed transcript	-	-	TSL:3

The strategy is based on the design of *Snrnp40-201* transcript, the transcription is shown below:



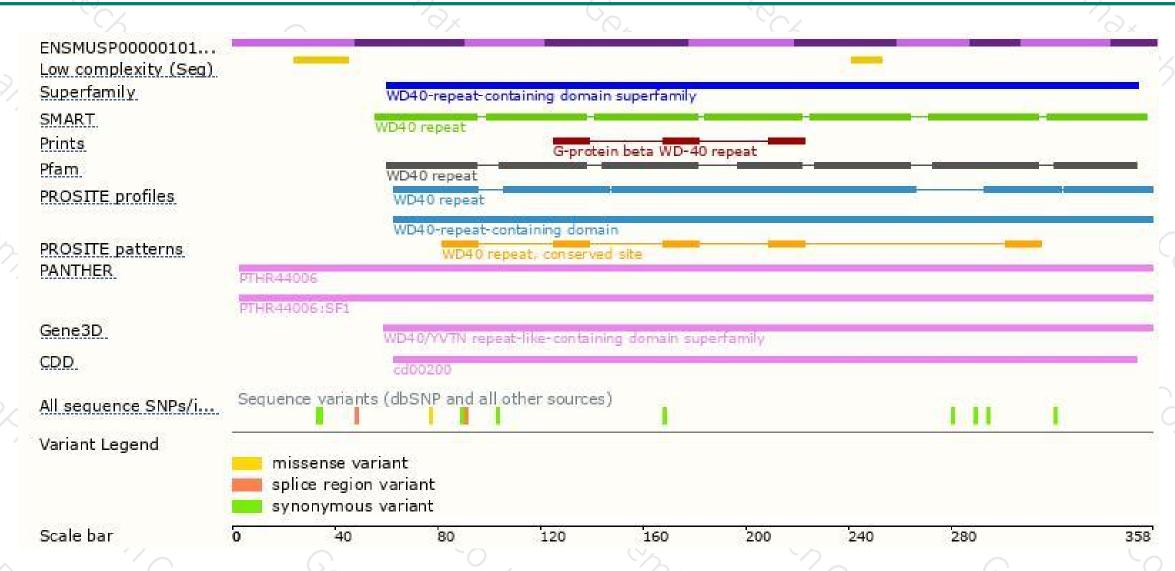
Genomic location distribution





Protein domain







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





