

Slc25a34 Cas9-KO Strategy

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Project Overview



Project Name Slc25a34

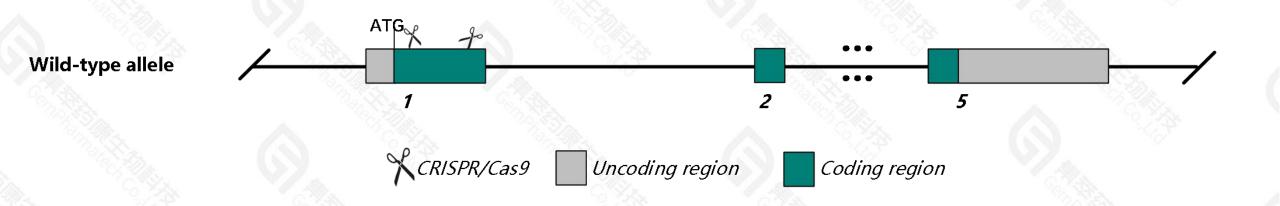
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Slc25a34 gene. The schematic diagram is as follows:



Technical routes



- > The Slc25a34 gene has 1 transcript. According to the structure of Slc25a34 gene, paritial coding sequence of exon1 of Slc25a34-201(ENSMUST00000038661.8) transcript is recommended as the knockout region. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Slc25a34* 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 with frameshift mutation of *Slc25a34* gene 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 *Slc25a34* 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)



Slc25a34 solute carrier family 25, member 34 [Mus musculus (house mouse)]

Gene ID: 384071, updated on 17-Dec-2020

Summary

☆ ?

Official Symbol Slc25a34 provided by MGI

Official Full Name solute carrier family 25, member 34 provided by MGI

Primary source MGI:MGI:2686215

See related Ensembl:ENSMUSG00000040740

RefSeq status PROVISIONAL
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as Gm1369

Expression Biased expression in heart adult (RPKM 69.7), colon adult (RPKM 20.1) and 5 other tissuesSee more

Orthologs <u>human all</u>

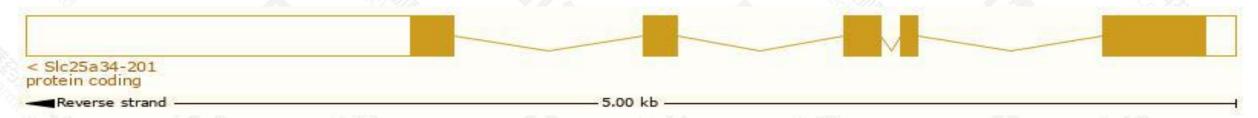
Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

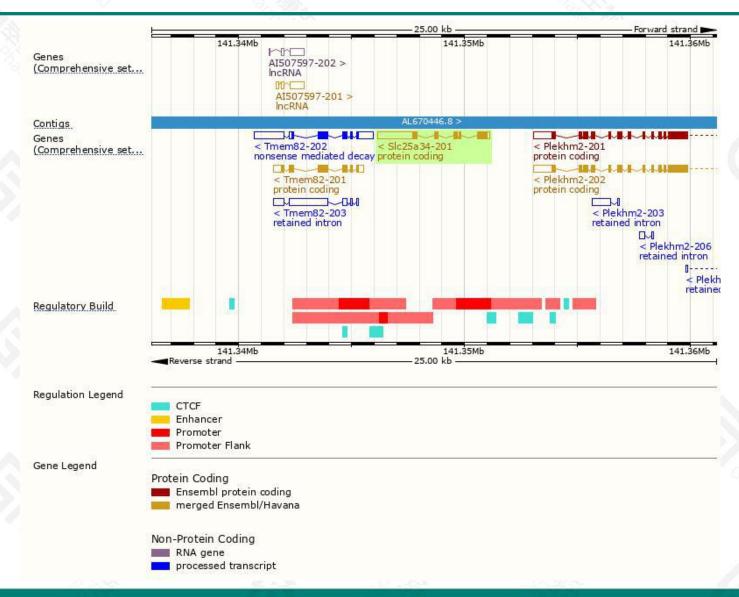
	Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
	Slc25a34-201	ENSMUST00000038661.8	2670	318aa	Protein coding	CCDS18878		TSL:1 , GENCODE basic , APPRIS P1 ,

The strategy is based on the design of *Slc25a34-201* transcript, the transcription is shown below:



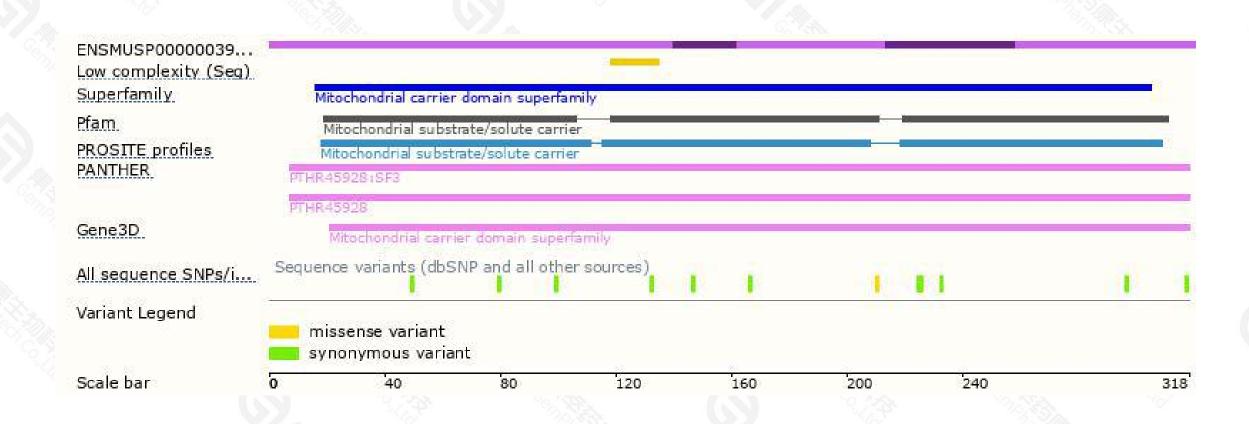
Genomic location distribution





Protein domain







If you have any questions, you are welcome to inquire.

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