

Slc22a16 Cas9-KO Strategy

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Project Overview

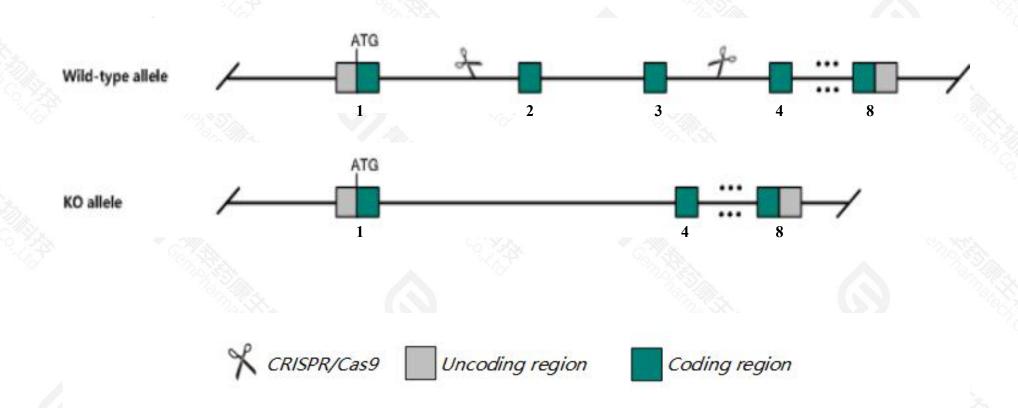


Project Name	Slc22a16			
Project type	Cas9-KO			
Strain background	C57BL/6JGpt			

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Slc22a16 gene. The schematic diagram is as follows:



Technical routes



- The *Slc22a16* gene has 2 transcripts. According to the structure of *Slc22a16* gene, exon2-exon3 of *Slc22a16*-202(ENSMUST00000078314.14) transcript is recommended as the knockout region. The region contains 598bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Slc22a16* 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 *Slc22a16* gene is located on the Chr10. 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)



Slc22a16 solute carrier family 22 (organic cation transporter), member 16 [Mus musculus (house mouse)]

Gene ID: 70840, updated on 17-Dec-2020

Summary



Official Symbol Slc22a16 provided by MGI

Official Full Name solute carrier family 22 (organic cation transporter), member 16 provided by MGI

Primary source MGI:MGI:1918090

See related Ensembl: ENSMUSG00000019834

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 4921504E14Rik, CT2, FLI, FLIPT2, OCT, OCT6, OKB, OKB1

Expression Restricted expression toward testis adult (RPKM 66.4)See more

Orthologs <u>human</u> all

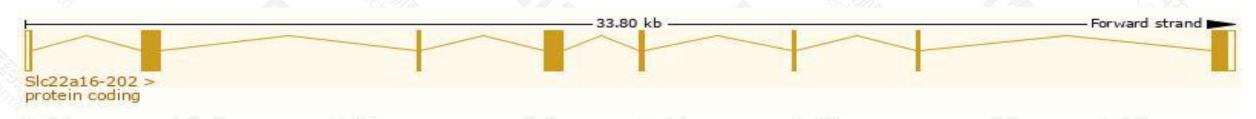
Transcript information (Ensembl)



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

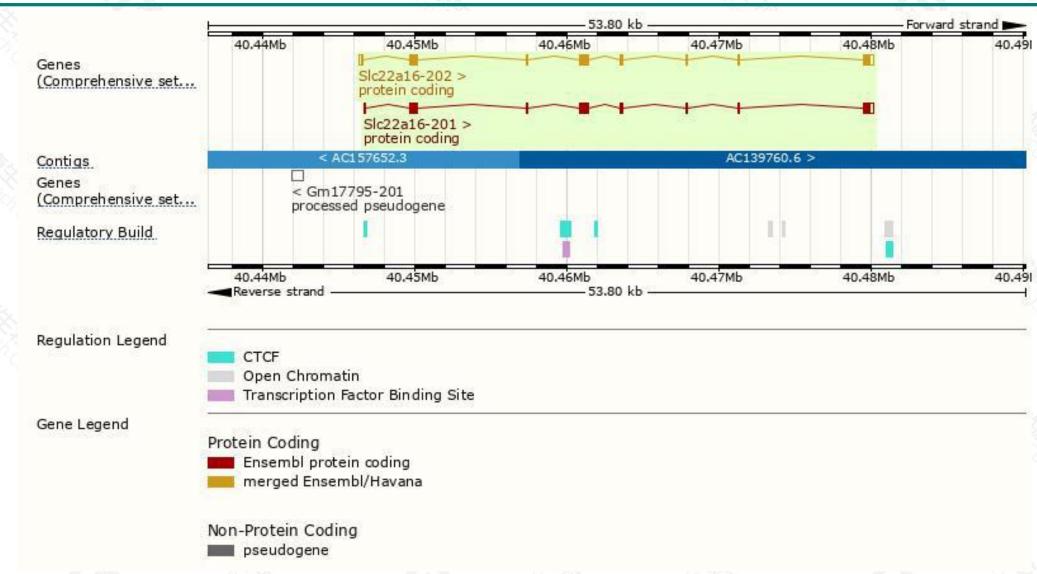
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc22a16-202	ENSMUST00000078314.14	2269	<u>649aa</u>	Protein coding	CCDS23796		TSL:1 , GENCODE basic , APPRIS P2 ,
Slc22a16-201	ENSMUST00000019978.9	2226	670aa	Protein coding	-		TSL:1, GENCODE basic, APPRIS ALT2,

The strategy is based on the design of *Slc22a16-202* transcript, the transcription is shown below:



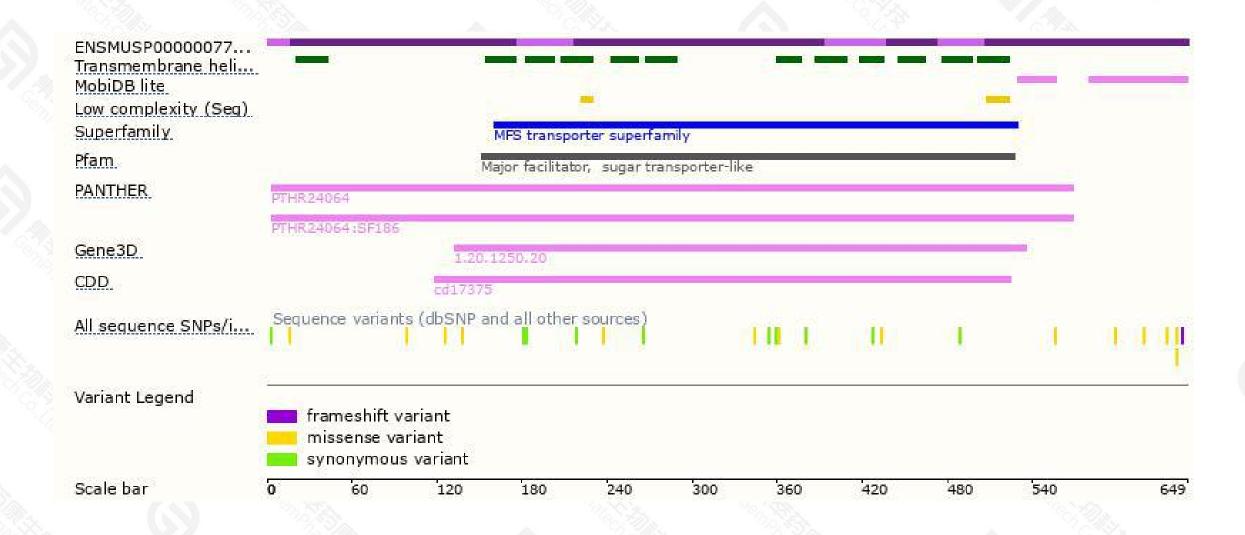
Genomic location distribution





Protein domain







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

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