

Slc31a1 Cas9-CKO Strategy

Designer:

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Design Date:

2019-8-15

Project Overview



Project Name

Slc31a1

Project type

Cas9-CKO

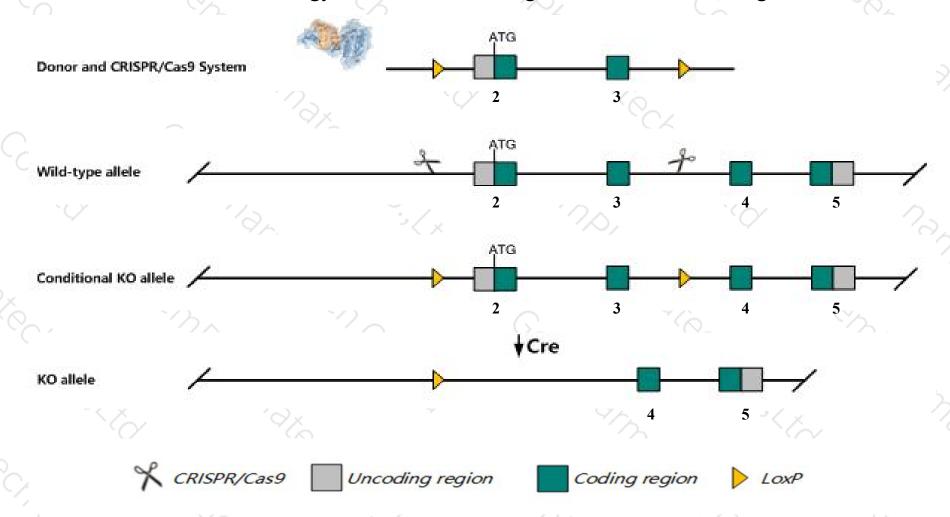
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Slc31a1* gene. The schematic diagram is as follows:



Technical routes



- The *Slc31a1* gene has 2 transcripts. According to the structure of *Slc31a1* gene, exon2-exon3 of *Slc31a1-201* (ENSMUST00000084526.11) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Slc31a1* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- > According to the existing MGI data, Mice homozygous for a null allele exhibit embryonic lethality during organogenesis associated with abnormal embryogenesis. Mice heterozygous for a null allele exhibit decreased copper levels in the blood and several organs.
- \succ The KO region contains functional region of the Cdc26-203. Knockout the region may affect the function of Cdc26-203.
- > The *Slc31a1* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



SIc31a1 solute carrier family 31, member 1 [Mus musculus (house mouse)]

Gene ID: 20529, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Slc31a1 provided by MGI

Official Full Name solute carrier family 31, member 1 provided by MGI

Primary source MGI:MGI:1333843

See related Ensembl:ENSMUSG00000066150

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 4930445G01Rik, Al787263, AU016967, Ctr1

Expression Ubiquitous expression in liver adult (RPKM 31.6), large intestine adult (RPKM 31.4) and 28 other tissuesSee more

Orthologs <u>human</u> all

Transcript information (Ensembl)



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

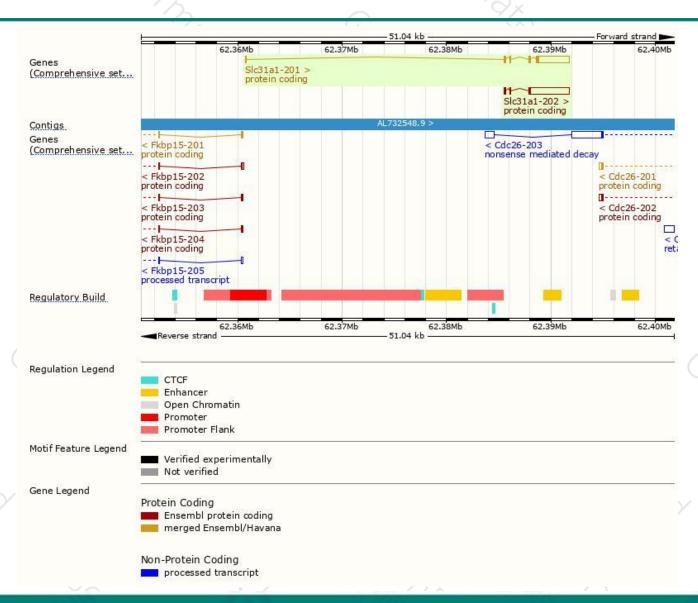
Name 🝦	Transcript ID	bp 🌲	Protein	Biotype 🖕	CCDS	UniProt 🍦	Flags
Slc31a1-201	ENSMUST00000084526.11	3718	<u>196aa</u>	Protein coding	CCDS18237 ₺	<u>Q8K211</u> ₽	TSL:1 GENCODE basic APPRIS P1
Slc31a1-202	ENSMUST00000122092.1	4154	<u>130aa</u>	Protein coding	384	<u>A8Y5P1</u> ₽	TSL:1 GENCODE basic

The strategy is based on the design of Slc31a1-201 transcript, The transcription is shown below

Slc31a1-201 > protein coding

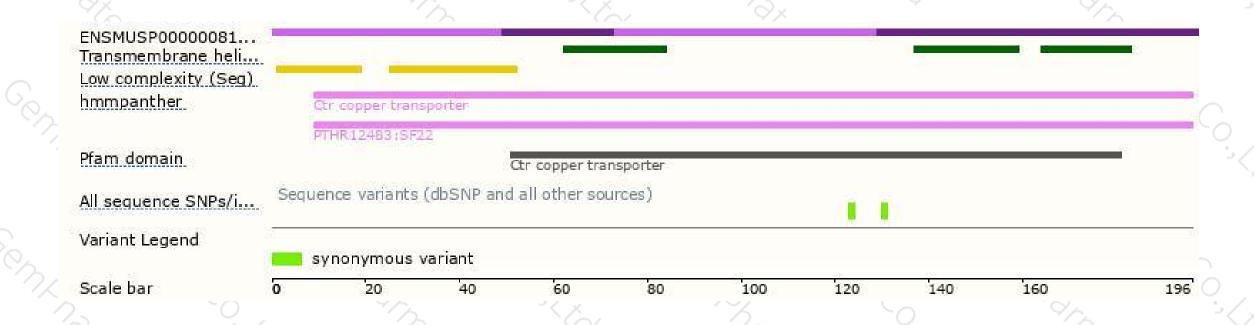
Genomic location distribution





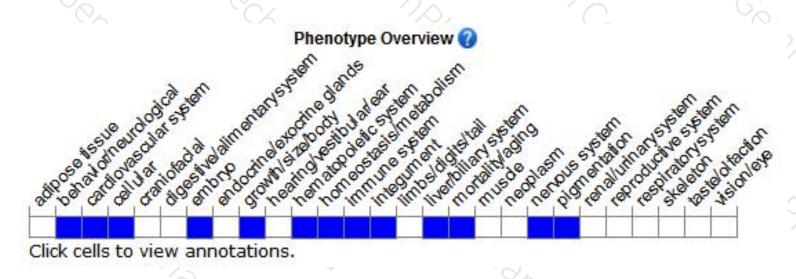
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

According to the existing MGI data, Mice homozygous for a null allele exhibit embryonic lethality during organogenesis associated with abnormal embryogenesis. Mice heterozygous for a null allele exhibit decreased copper levels in the blood and several organs.



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





