

Slc17a9 Cas9-CKO Strategy

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Project Overview



Project Name

Slc17a9

Project type

Cas9-CKO

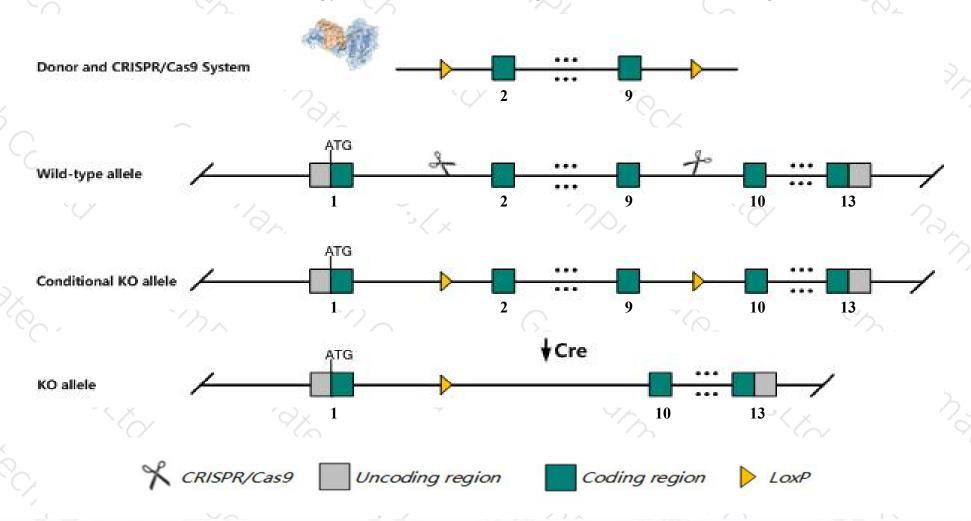
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Slc17a9 gene. The schematic diagram is as follows:



Technical routes



- ➤ The Slc17a9 gene has 5 transcripts. According to the structure of Slc17a9 gene, exon2-exon9 of Slc17a9-201 (ENSMUST00000094218.3) transcript is recommended as the knockout region. The region contains 886bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc17a9* 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, Homozygous knockout affects the neuroendocrine system, resulting in hypoglycemia, increased glucose tolerance and increased insulin sensitivity.
- > The Slc17a9 gene is located on the Chr2. 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)



SIc17a9 solute carrier family 17, member 9 [Mus musculus (house mouse)]

Gene ID: 228993, updated on 6-Apr-2019

Summary

↑ ?

Official Symbol Slc17a9 provided by MGI

Official Full Name solute carrier family 17, member 9 provided by MGI

Primary source MGI:MGI:1919107

See related Ensembl:ENSMUSG00000023393

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 1700019H03Rik, BC019537, Vnut

Expression Biased expression in colon adult (RPKM 55.9), genital fat pad adult (RPKM 48.4) and 13 other tissuesSee more

Orthologs <u>human</u> all

Transcript information (Ensembl)



The gene has 5 transcripts, all transcripts are shown below:

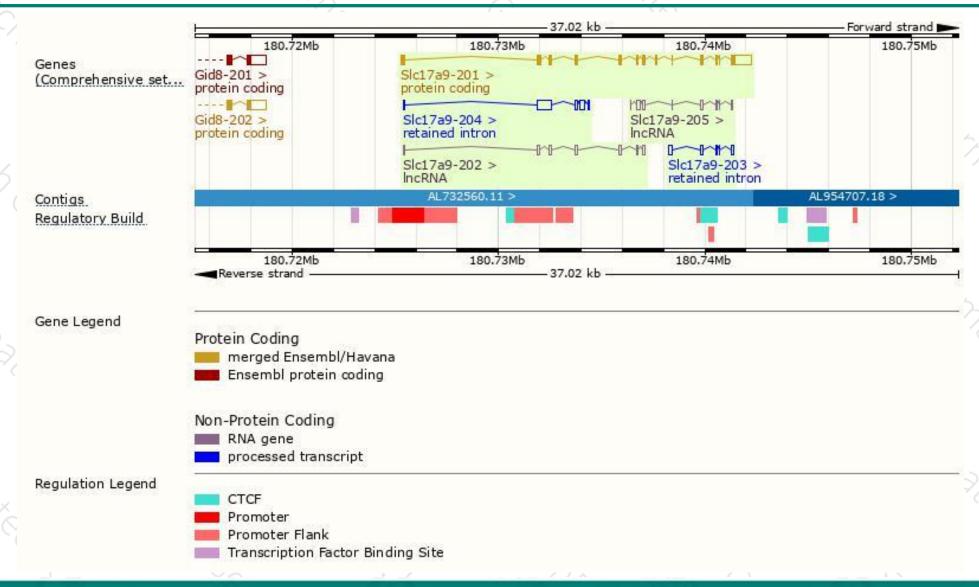
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
SIc17a9-201	ENSMUST00000094218.3	2275	<u>447aa</u>	Protein coding	CCDS38378	Q8VCL5	TSL:5 GENCODE basic APPRIS P1
SIc17a9-204	ENSMUST00000151882.7	1158	No protein	Retained intron	9 4 3	-	TSL:5
SIc17a9-203	ENSMUST00000140955.1	506	No protein	Retained intron	820	-	TSL:3
Slc17a9-202	ENSMUST00000139817.1	754	No protein	IncRNA	1528	-	TSL:5
SIc17a9-205	ENSMUST00000154882.7	621	No protein	IncRNA	-		TSL:1

The strategy is based on the design of Slc17a9-201 transcript, The transcription is shown below



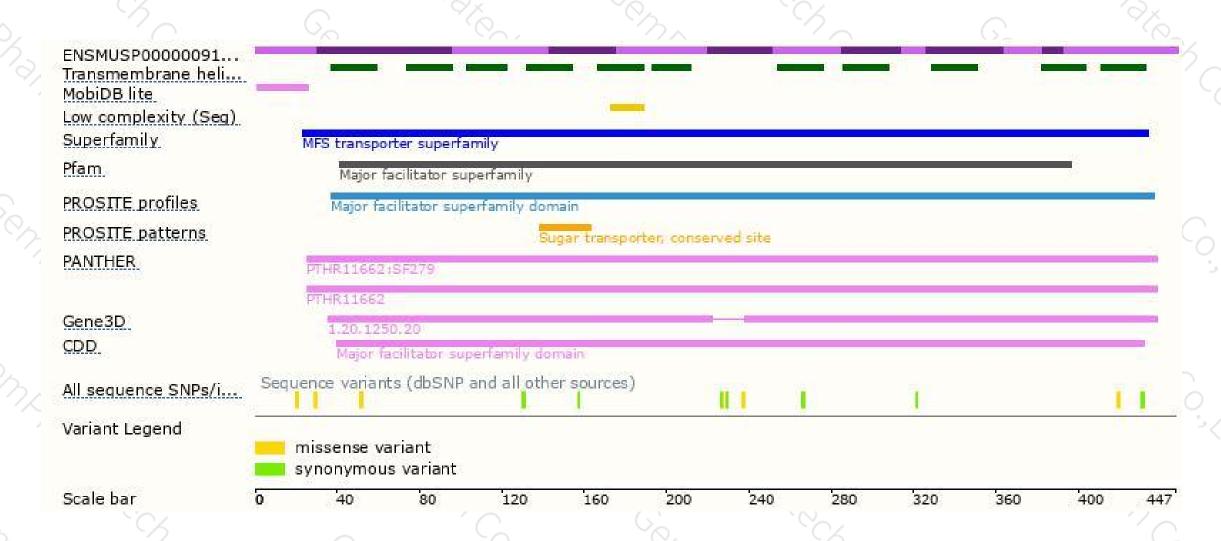
Genomic location distribution





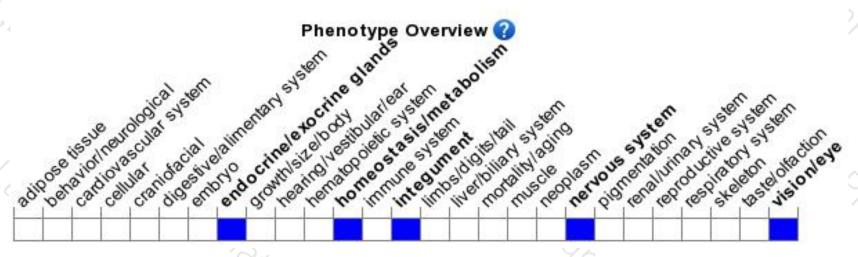
Protein domain





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, Homozygous knockout affects the neuroendocrine system, resulting in hypoglycemia, increased glucose tolerance and increased insulin sensitivity.



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





