

Slc13a3 Cas9-CKO Strategy

Designer: Rui Xiong

Reviewer: Miaomiao Cui

Design Date: 2023-06-01

Overview

Target Gene Name

• Slc13a3

Project Type

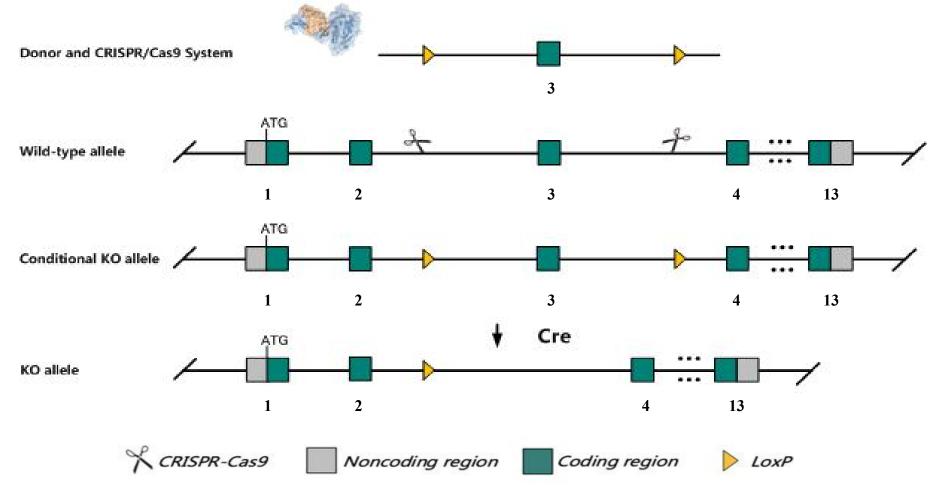
• Cas9-CKO

Genetic Background

• C57BL/6JGpt



Strain Strategy



Schematic representation of CRISPR-Cas9 engineering used to edit the Slc13a3 gene.



Technical Information

- The *Slc13a3* gene has 5 transcripts. According to the structure of *Slc13a3* gene, exon3 of *Slc13a3*-201 (ENSMUST00000029208.15) transcript is recommended as the knockout region. The region contains 164bp coding sequence. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Slc13a3* 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 on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.
- 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.



Gene Information

SIc13a3 solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3 [Mus musculus (house mouse)]

Gene ID: 114644, updated on 31-Jan-2019



Source: https://www.ncbi.nlm.nih.gov/

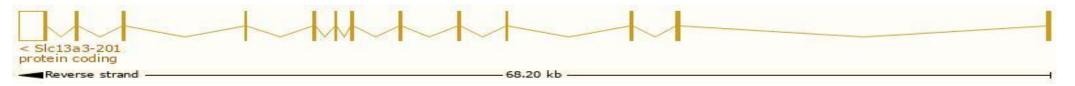


Transcript Information

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
SIc13a3-201	ENSMUST00000029208.14	3524	<u>600aa</u>	Protein coding	CCDS17081	Q3UUJ6 Q91Y63	TSL:1 GENCODE basic APPRIS P2
SIc13a3-202	ENSMUST00000109279.2	3131	<u>558aa</u>	Protein coding	-	<u>A2A4U9</u>	TSL:1 GENCODE basic APPRIS ALT2
SIc13a3-205	ENSMUST00000147107.1	1968	No protein	Retained intron	XI-	12	TSL:1
SIc13a3-203	ENSMUST00000143136.1	456	No protein	Retained intron	-	725	TSL:2
SIc13a3-204	ENSMUST00000145513.1	397	No protein	Retained intron		(2)	TSL:2

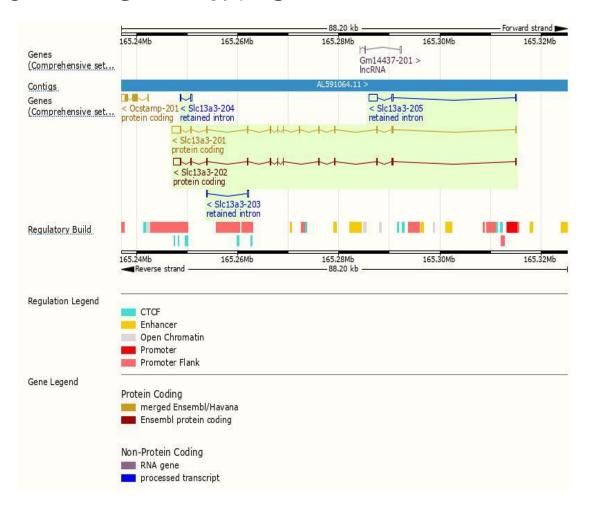
The strategy is based on the design of *Slc13a3*-201 transcript, the transcription is shown below:



Source: https://www.ensembl.org



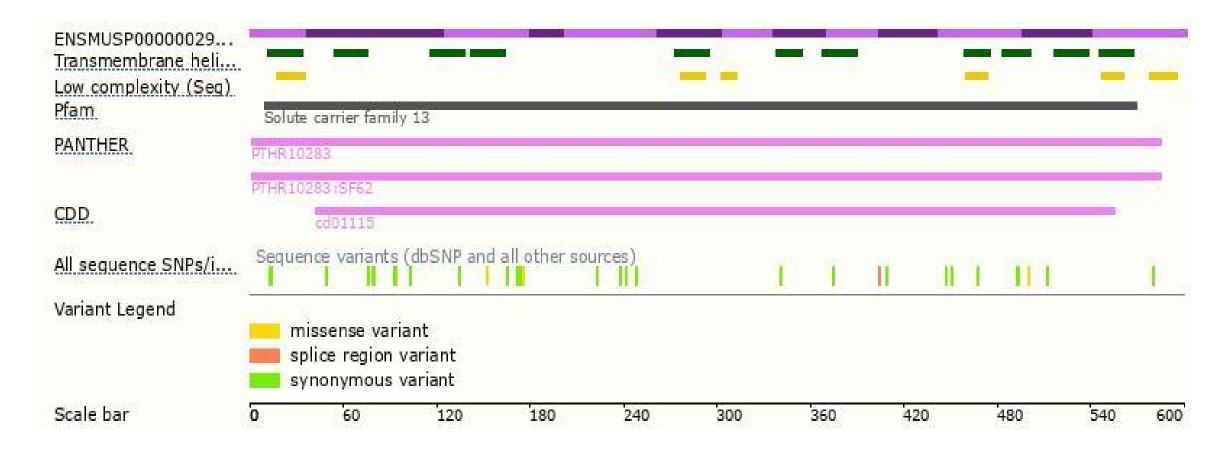
Genomic Information

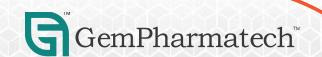




Source: : https://www.ensembl.org

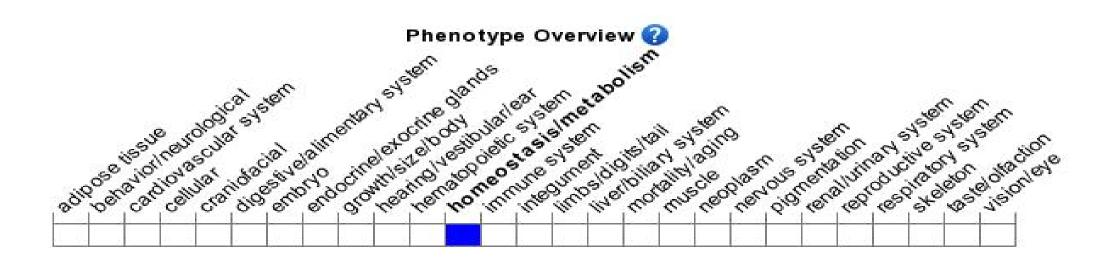
Protein Information





Source: : https://www.ensembl.org

Mouse Phenotype Information (MGI)



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



Source: https://www.informatics.jax.org

Important Information

- The flox region contains functional region of the Gm14437-201 gene. Knockout the region may affect the function of Gm14437-201 gene.
- *Slc13a3* is located on Chr2. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is 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 the existing technology level.

