

Slc38a9 Cas9-KO Strategy

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Project Overview

Project Name

Slc38a9

Project type

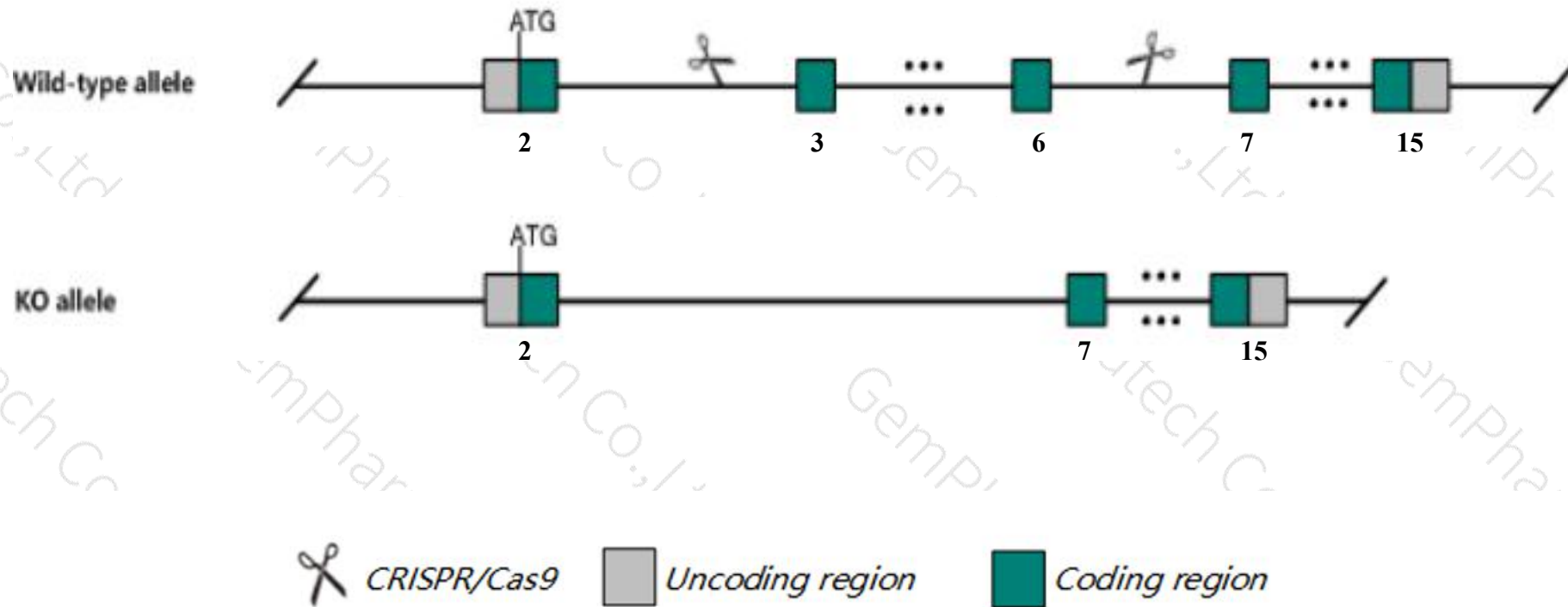
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc38a9* gene. The schematic diagram is as follows:



- The *Slc38a9* gene has 8 transcripts. According to the structure of *Slc38a9* gene, exon3-exon6 of *Slc38a9-201* (ENSMUST00000052514.5) transcript is recommended as the knockout region. The region contains 413bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc38a9* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Slc38a9* gene is located on the Chr13. 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.
- The function of *Gm47827* gene may be affect.
- 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)

Slc38a9 solute carrier family 38, member 9 [*Mus musculus* (house mouse)]

Gene ID: 268706, updated on 13-Mar-2020

Summary

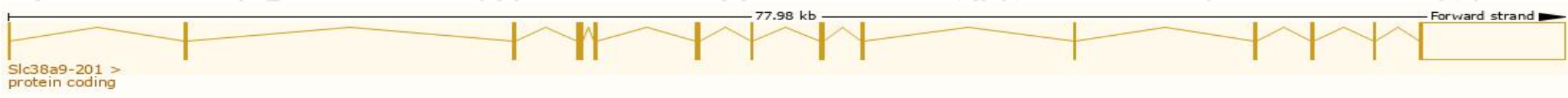
Official Symbol	Slc38a9 provided by MGI
Official Full Name	solute carrier family 38, member 9 provided by MGI
Primary source	MGI:MGI:1918839
See related	Ensembl:ENSMUSG00000047789
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	A730092C09; 4833412L08Rik; 6720411P22Rik; 9130023D20Rik; 9430067K09Rik
Expression	Ubiquitous expression in testis adult (RPKM 7.7), thymus adult (RPKM 4.8) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

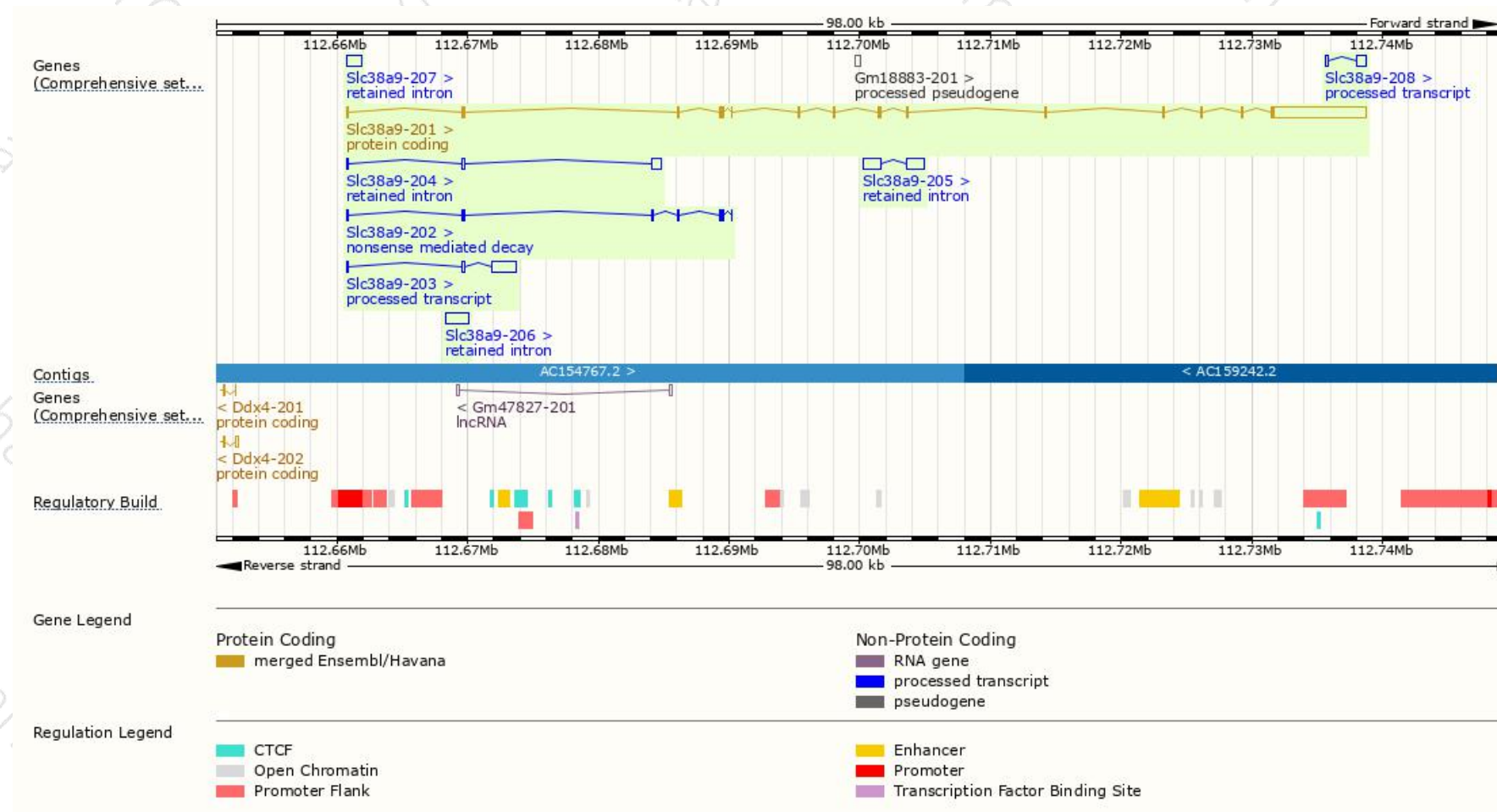
The gene has 8 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc38a9-201	ENSMUST00000052514.5	8910	560aa	Protein coding	CCDS26775	Q8BGD6	TSL:1 GENCODE basic APPRIS P1
Slc38a9-202	ENSMUST00000223581.1	681	42aa	Nonsense mediated decay	-	A0A286YCC6	-
Slc38a9-203	ENSMUST00000223674.1	2063	No protein	Processed transcript	-	-	-
Slc38a9-208	ENSMUST00000225649.1	1011	No protein	Processed transcript	-	-	-
Slc38a9-205	ENSMUST00000224252.1	2847	No protein	Retained intron	-	-	-
Slc38a9-206	ENSMUST00000224703.1	1768	No protein	Retained intron	-	-	-
Slc38a9-207	ENSMUST00000225367.1	1178	No protein	Retained intron	-	-	-
Slc38a9-204	ENSMUST00000223839.1	980	No protein	Retained intron	-	-	-

The strategy is based on the design of *Slc38a9-201* 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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