

Slc5a11 Cas9-KO Strategy

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Project Overview

Project Name

Slc5a11

Project type

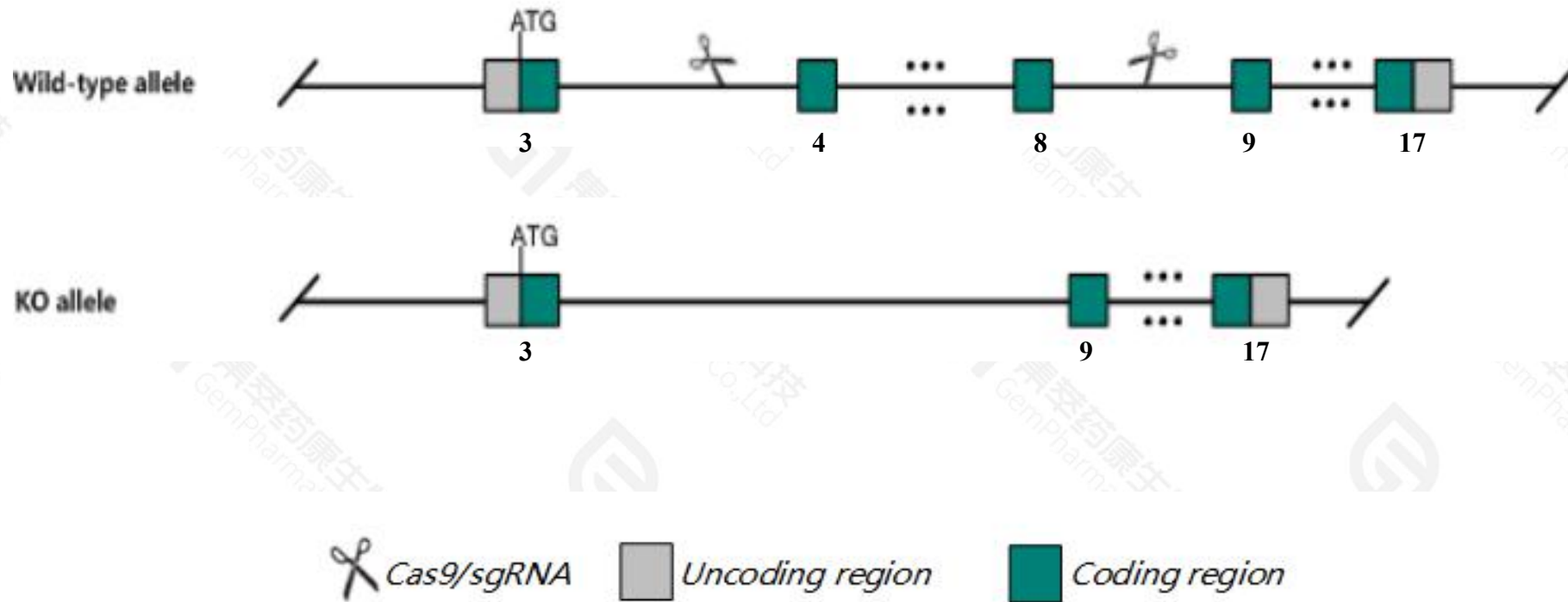
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc5a11* gene. The schematic diagram is as follows:



- The *Slc5a11* gene has 8 transcripts. According to the structure of *Slc5a11* gene, exon4-exon8 of *Slc5a11-201*(ENSMUST00000033035.13) transcript is recommended as the knockout region. The region contains 448bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc5a11* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA 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 *Slc5a11* gene is located on the Chr7. 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.

Slc5a11 solute carrier family 5 (sodium/glucose cotransporter), member 11 [Mus musculus (house mouse)]

Gene ID: 233836, updated on 10-Oct-2020

Summary



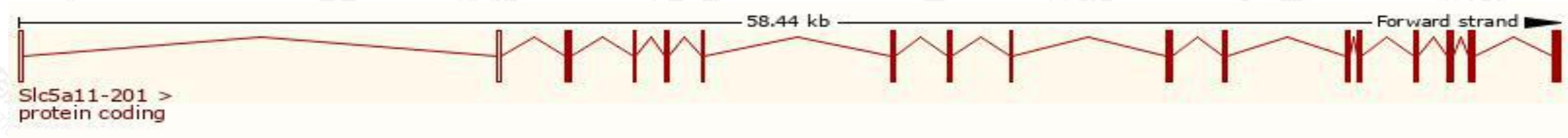
| | |
|---------------------------|---|
| Official Symbol | Slc5a11 provided by MGI |
| Official Full Name | solute carrier family 5 (sodium/glucose cotransporter), member 11 provided by MGI |
| Primary source | MGI:MGI:1919316 |
| See related | Ensembl:ENSMUSG00000030769 |
| Gene type | protein coding |
| RefSeq status | PROVISIONAL |
| Organism | Mus musculus |
| Lineage | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| Also known as | 2010013B02Rik, Kst, Kst1, RKST2, SMIT2, Slc5a10 |
| Expression | Biased expression in duodenum adult (RPKM 46.1), small intestine adult (RPKM 29.5) and 2 other tissues See more |
| Orthologs | human all |

Transcript information (Ensembl)

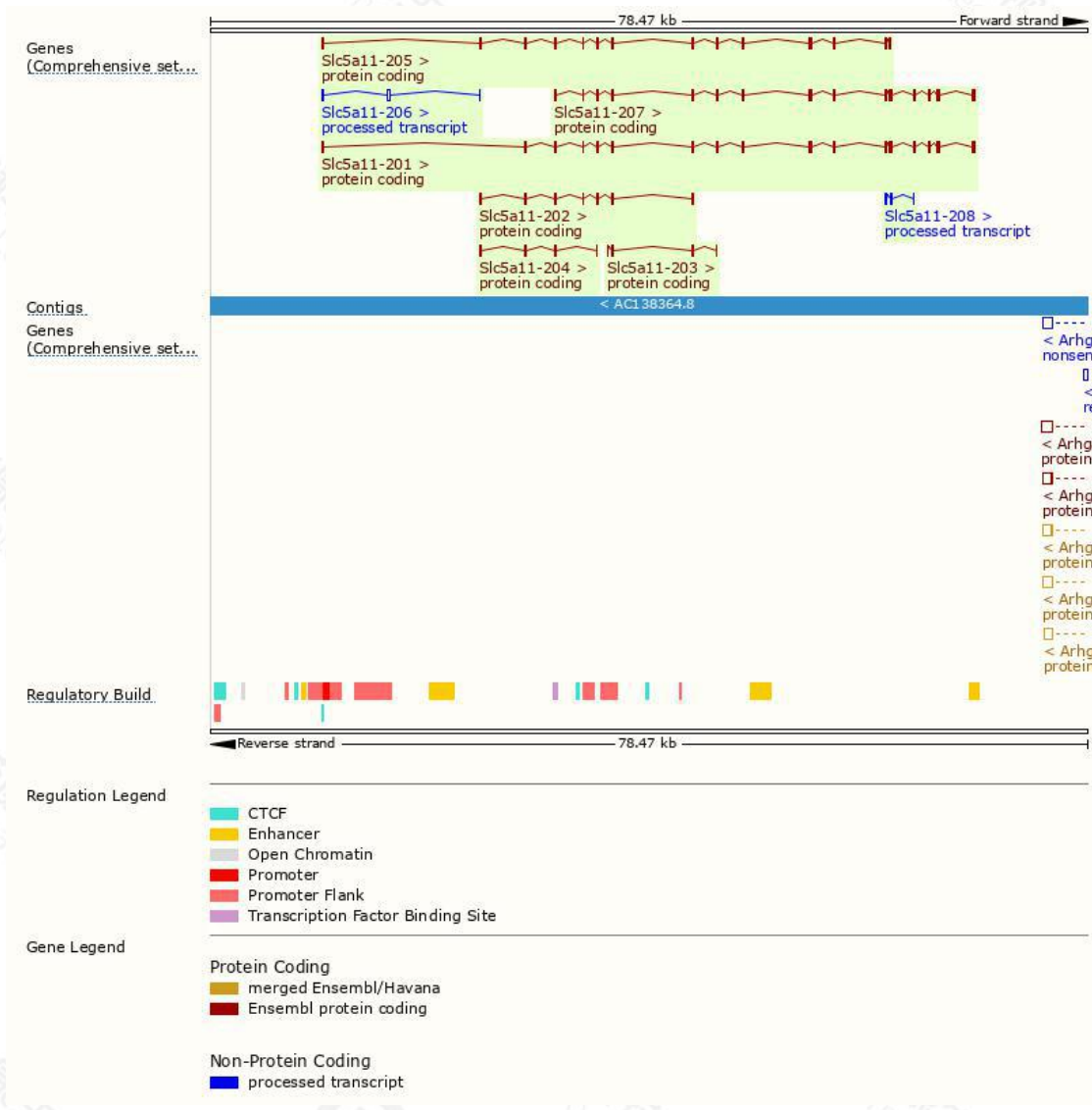
The gene has 8 transcripts,all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|-------------|---------------------------------------|------|-----------------------|----------------------|---------------------------|---------|-------------------------------------|
| Slc5a11-201 | ENSMUST00000033035.13 | 2396 | 673aa | Protein coding | CCDS21819 | | TSL:1 , GENCODE basic , APPRIS P1 , |
| Slc5a11-207 | ENSMUST00000167299.9 | 2022 | 673aa | Protein coding | CCDS21819 | | TSL:5 , GENCODE basic , APPRIS P1 , |
| Slc5a11-205 | ENSMUST00000131933.8 | 1648 | 402aa | Protein coding | - | | CDS 3' incomplete , TSL:5 , |
| Slc5a11-202 | ENSMUST00000127655.8 | 704 | 152aa | Protein coding | - | | CDS 3' incomplete , TSL:3 , |
| Slc5a11-204 | ENSMUST00000131461.3 | 440 | 64aa | Protein coding | - | | CDS 3' incomplete , TSL:3 , |
| Slc5a11-203 | ENSMUST00000131209.2 | 215 | 72aa | Protein coding | - | | CDS 5' and 3' incomplete , TSL:5 , |
| Slc5a11-206 | ENSMUST00000140721.2 | 459 | No protein | Processed transcript | - | | TSL:5 , |
| Slc5a11-208 | ENSMUST00000206180.2 | 242 | No protein | Processed transcript | - | | TSL:5 , |

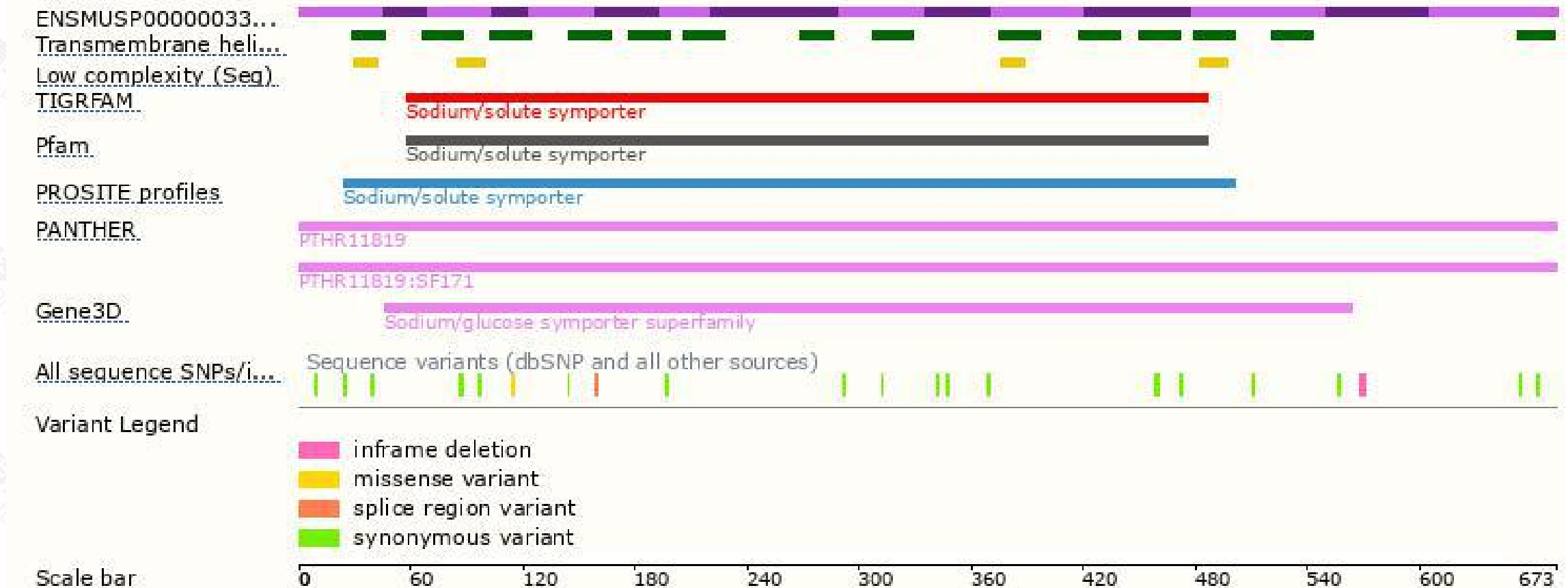
The strategy is based on the design of *Slc5a11-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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