

# ***Slc25a5*** Cas9-KO Strategy

Designer: Lixin Lv

# Project Overview

**Project Name**

*Slc25a5*

**Project type**

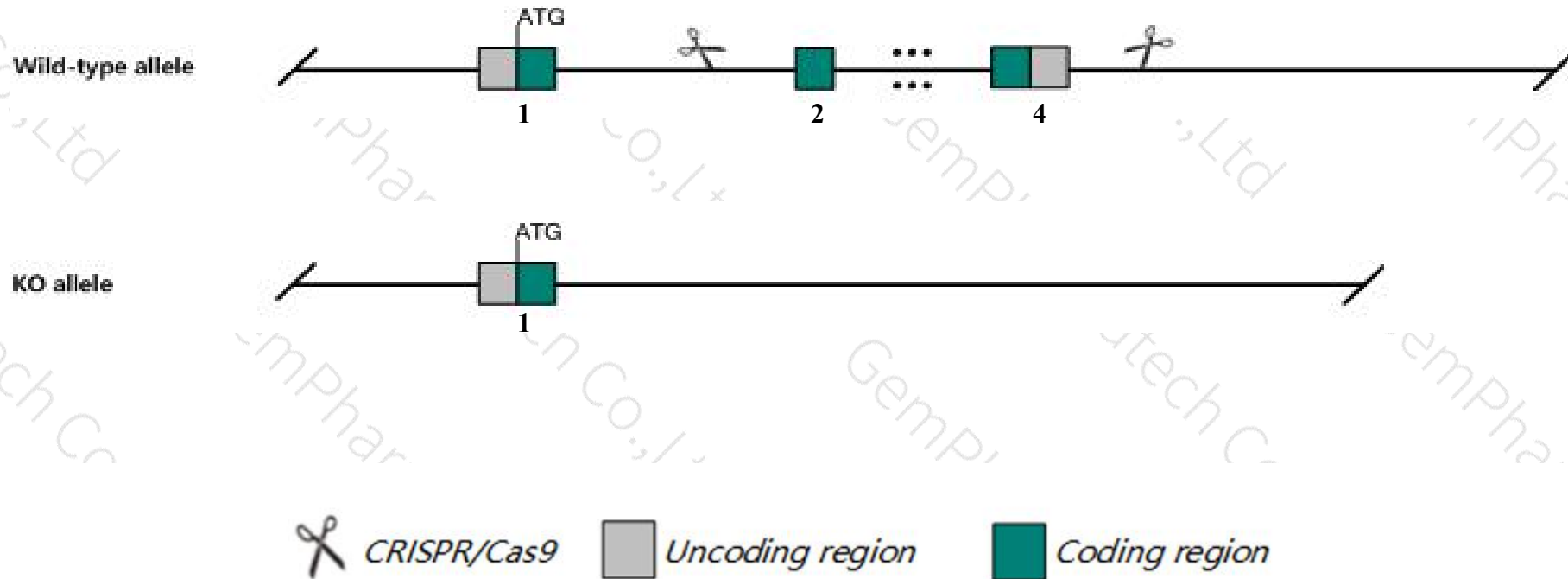
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc25a5* gene. The schematic diagram is as follows:



- The *Slc25a5* gene has 3 transcripts. According to the structure of *Slc25a5* gene, exon2-exon4 of *Slc25a5-201* (ENSMUST00000016463.3) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc25a5* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Slc25a5* gene is located on the ChrX. 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.



# Gene information (NCBI)

## Slc25a5 solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5 [Mus musculus (house mouse)]

Gene ID: 11740, updated on 7-Apr-2019

### Summary



**Official Symbol** Slc25a5 provided by [MGI](#)

**Official Full Name** solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5 provided by [MGI](#)

**Primary source** [MGI:MGI:1353496](#)

**See related** [Ensembl:ENSMUSG00000016319](#)

**Gene type** protein coding

**RefSeq status** REVIEWED

**Organism** [Mus musculus](#)

**Lineage** Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

**Also known as** Ant2

**Summary** This gene encodes a transmembrane domain-containing protein that localizes to the mitochondrial inner membrane. The encoded protein facilitates the exchange of ADP from the cytoplasm with ATP from the mitochondria. Pseudogenes for this gene are found on multiple chromosomes. [provided by RefSeq, May 2015]

**Expression** Ubiquitous expression in large intestine adult (RPKM 861.5), kidney adult (RPKM 667.1) and 27 other tissues [See more](#)

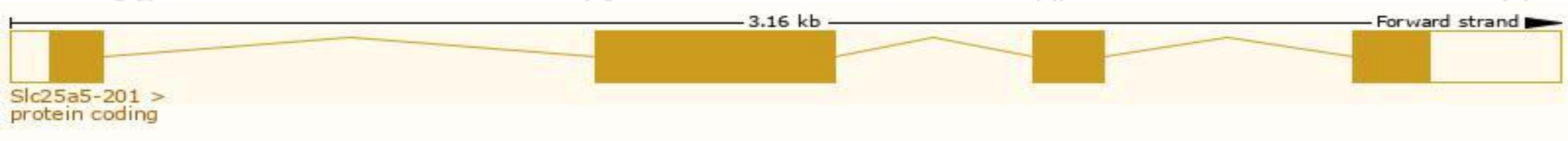
**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

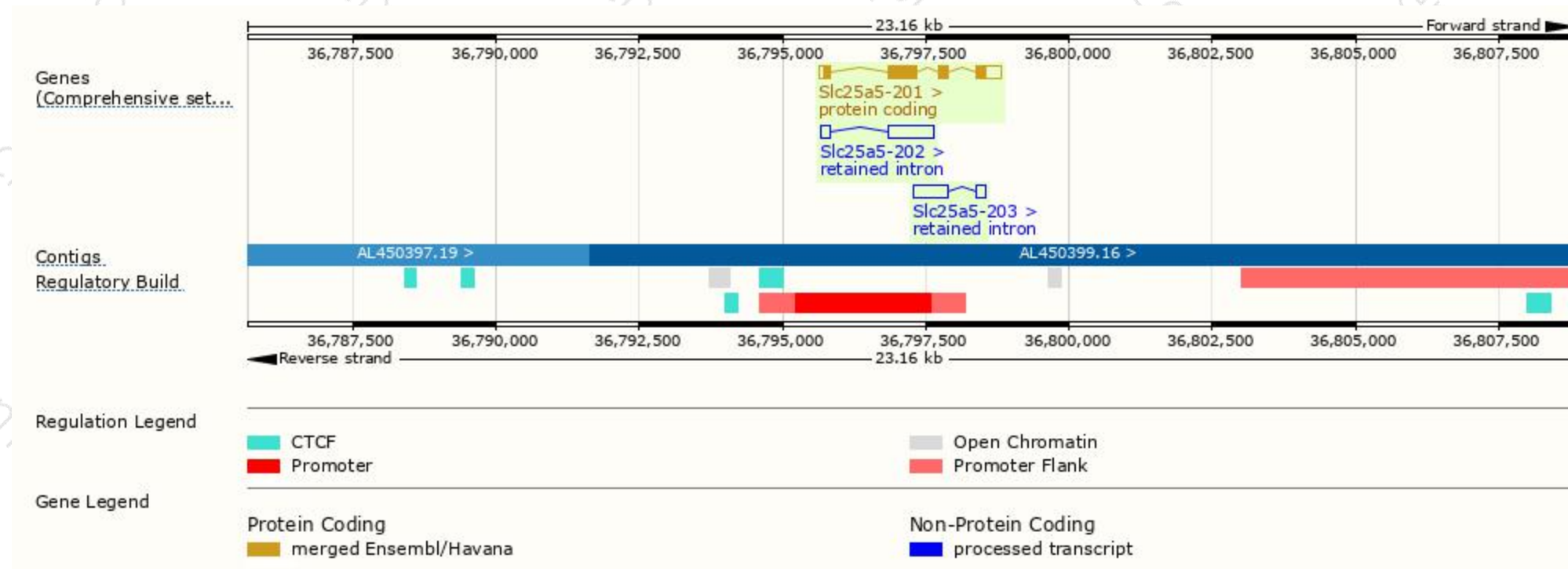
The gene has 3 transcripts,all transcripts are shown below:

| Name        | Transcript ID                        | bp   | Protein               | Biotype         | CCDS                      | UniProt                       | Flags                         |
|-------------|--------------------------------------|------|-----------------------|-----------------|---------------------------|-------------------------------|-------------------------------|
| Slc25a5-201 | <a href="#">ENSMUST00000016463.3</a> | 1240 | <a href="#">298aa</a> | Protein coding  | <a href="#">CCDS30062</a> | <a href="#">P51881 Q545A2</a> | TSL:1 GENCODE basic APPRIS P1 |
| Slc25a5-202 | <a href="#">ENSMUST00000126608.1</a> | 989  | No protein            | Retained intron | -                         | -                             | TSL:1                         |
| Slc25a5-203 | <a href="#">ENSMUST00000156672.1</a> | 765  | No protein            | Retained intron | -                         | -                             | TSL:1                         |

The strategy is based on the design of *Slc25a5-201* transcript,The transcription is shown below

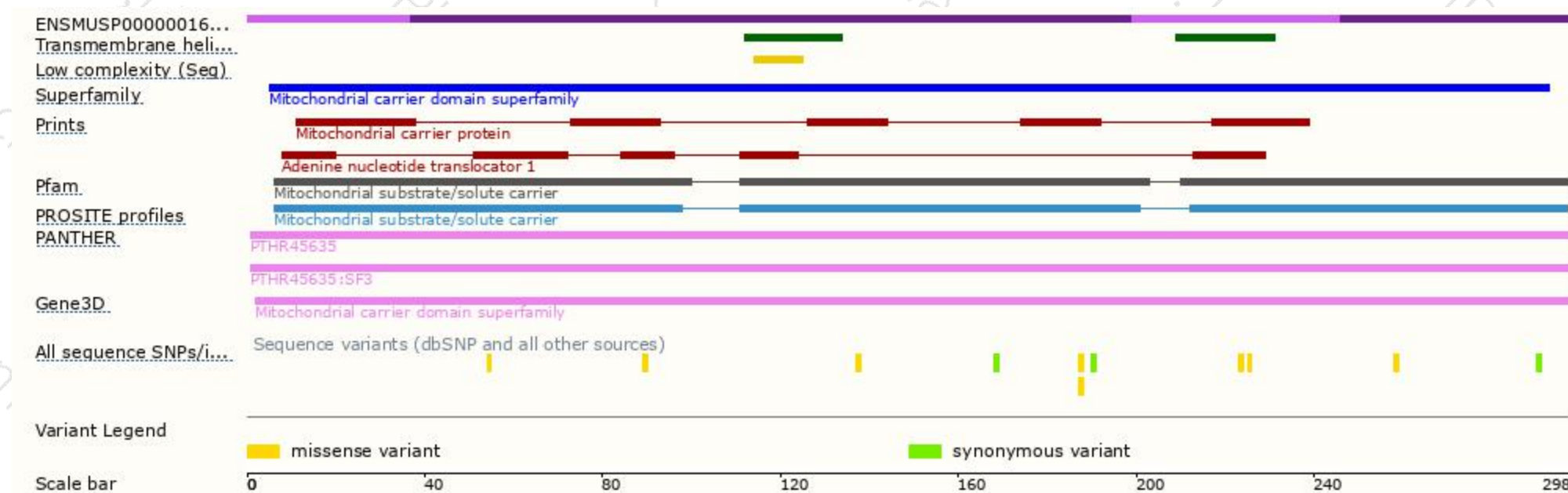


# Genomic location distribution





# Protein domain



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

Tel: 400-9660890

