

Tm9sf2 Cas9-KO Strategy

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Project Overview

Project Name

Tm9sf2

Project type

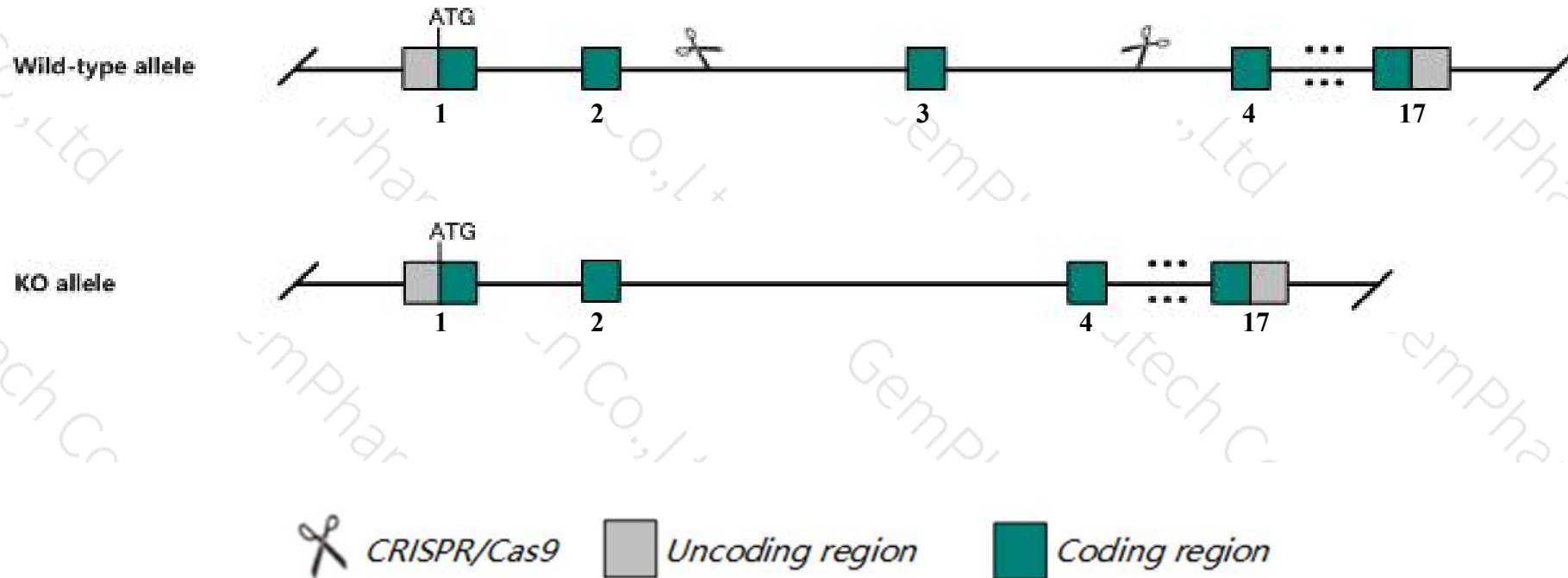
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Tm9sf2* gene. The schematic diagram is as follows:



- The *Tm9sf2* gene has 3 transcripts. According to the structure of *Tm9sf2* gene, exon3 of *Tm9sf2-201* (ENSMUST00000026624.10) transcript is recommended as the knockout region. The region contains 94bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Tm9sf2* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Tm9sf2* gene is located on the Chr14. 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)

Tm9sf2 transmembrane 9 superfamily member 2 [Mus musculus (house mouse)]

Gene ID: 68059, updated on 31-Jan-2019

Summary



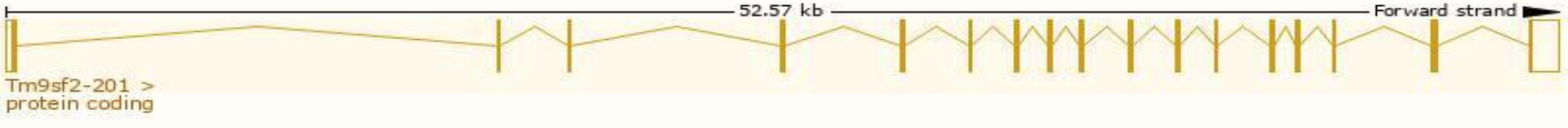
Official Symbol	Tm9sf2 provided by MGI
Official Full Name	transmembrane 9 superfamily member 2 provided by MGI
Primary source	MGI:MGI:1915309
See related	Ensembl:ENSMUSG00000025544
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1500001N15Rik, AA536814, D14Etd64e, P76
Expression	Ubiquitous expression in colon adult (RPKM 91.0), duodenum adult (RPKM 80.6) and 28 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
Tm9sf2-201	ENSMUST00000026624.10	3107	662aa	Protein coding	CCDS27345	P58021 Q542E4	TSL:1 GENCODE basic APPRIS P1
Tm9sf2-203	ENSMUST00000171318.1	1494	497aa	Protein coding	-	E9PZ69	TSL:5 GENCODE basic
Tm9sf2-202	ENSMUST00000170007.1	627	57aa	Nonsense mediated decay	-	E9Q850	TSL:3

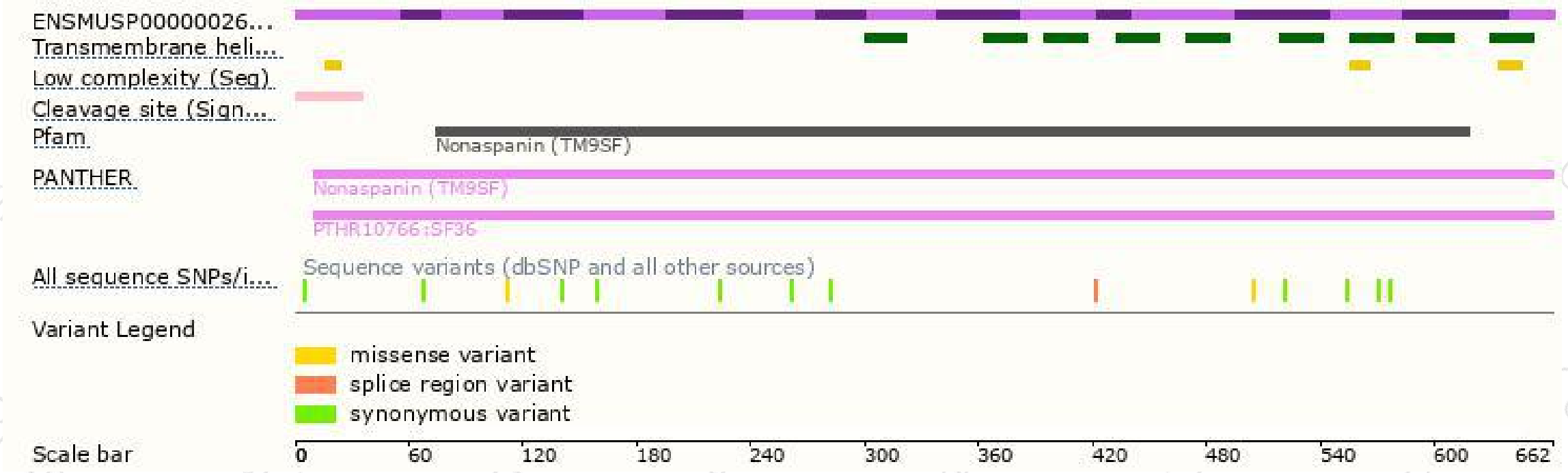
The strategy is based on the design of *Tm9sf2-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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