

Igsf6 Cas9-KO Strategy

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Project Overview

Project Name

Igsf6

Project type

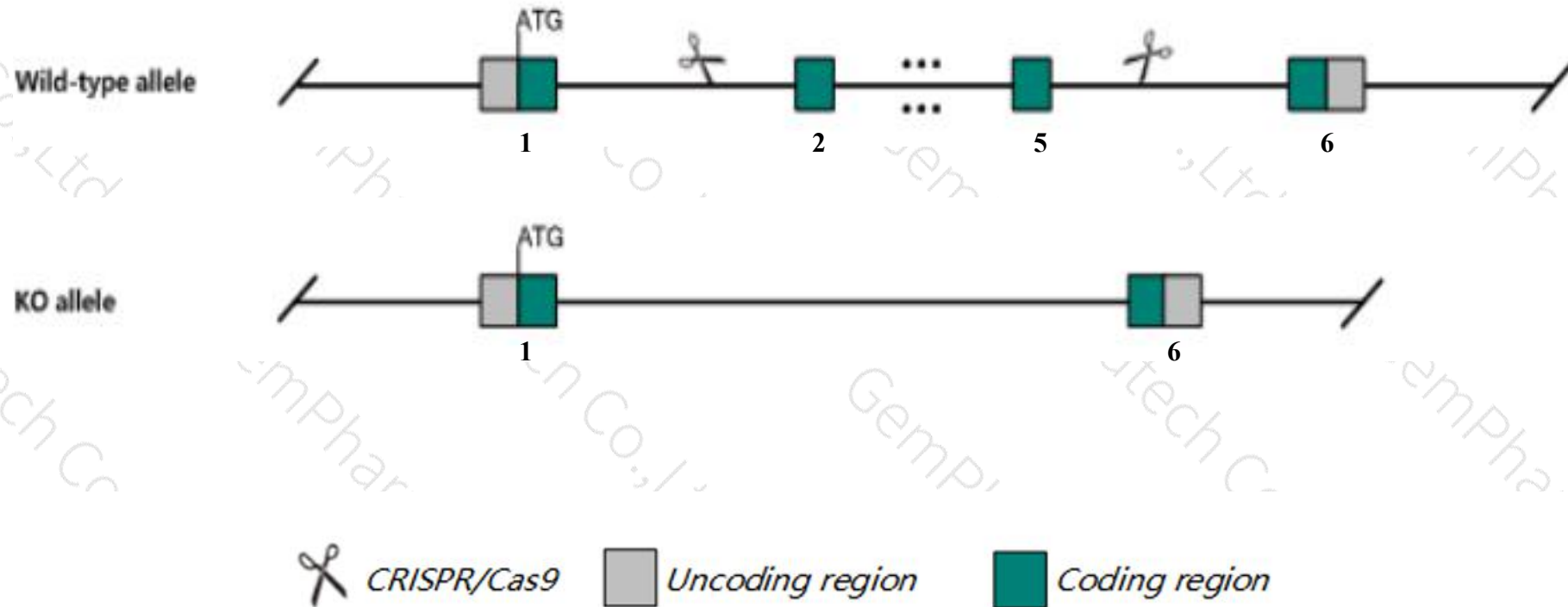
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Igsf6* gene. The schematic diagram is as follows:



- The *Igsf6* gene has 1 transcript. According to the structure of *Igsf6* gene, exon2-exon5 of *Igsf6*-201(ENSMUST00000047194.3) transcript is recommended as the knockout region. The region contains 590bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Igsf6* gene. The brief process is as follows: CRISPR/Cas9 system 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 knockout region is near to the N-terminal of *2210406H18Rik* gene, this strategy may influence the regulatory function of the N-terminal of *2210406H18Rik* gene.
- The partial intron of *Mettl9* gene will be deleted together in this strategy.
- The *Igsf6* 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.

Gene information (NCBI)

Igsf6 immunoglobulin superfamily, member 6 [Mus musculus (house mouse)]

Gene ID: 80719, updated on 13-Mar-2020

Summary



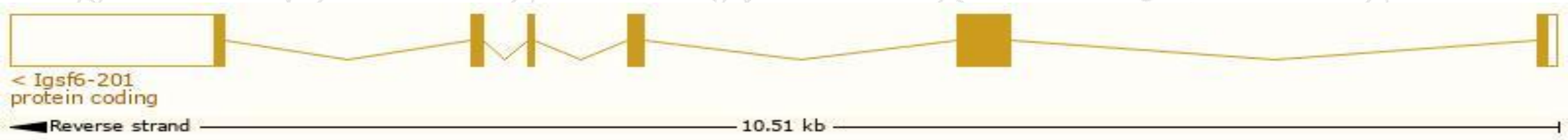
Official Symbol	Igsf6 provided by MGI
Official Full Name	immunoglobulin superfamily, member 6 provided by MGI
Primary source	MGI:MGI:1891393
See related	Ensembl:ENSMUSG00000035004
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Broad expression in liver E18 (RPKM 6.0), liver E14 (RPKM 3.2) and 19 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

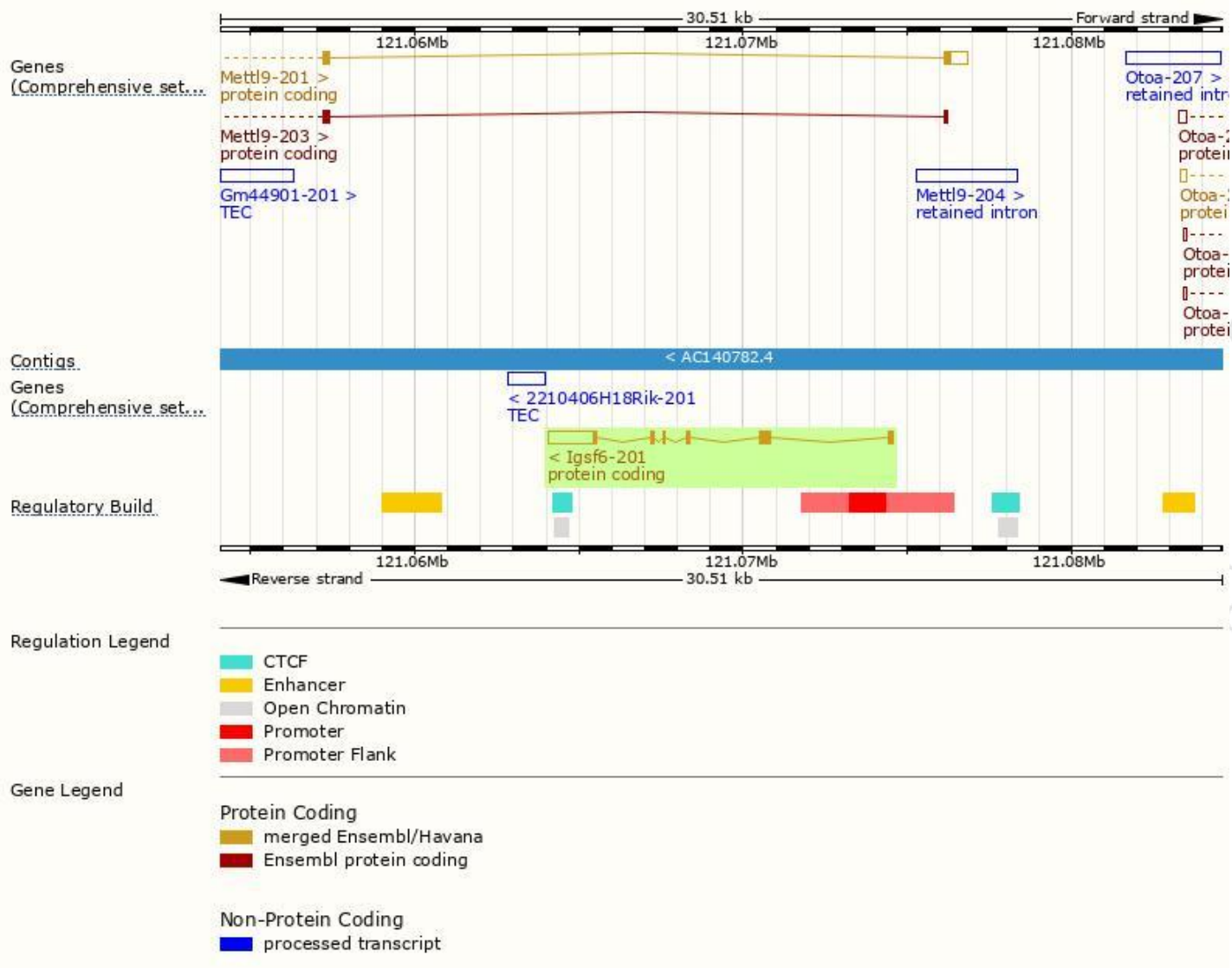
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Igsf6-201	ENSMUST00000047194.3	2175	237aa	Protein coding	CCDS21801	P0C6B7	TSL:1 GENCODE basic APPRIS P1

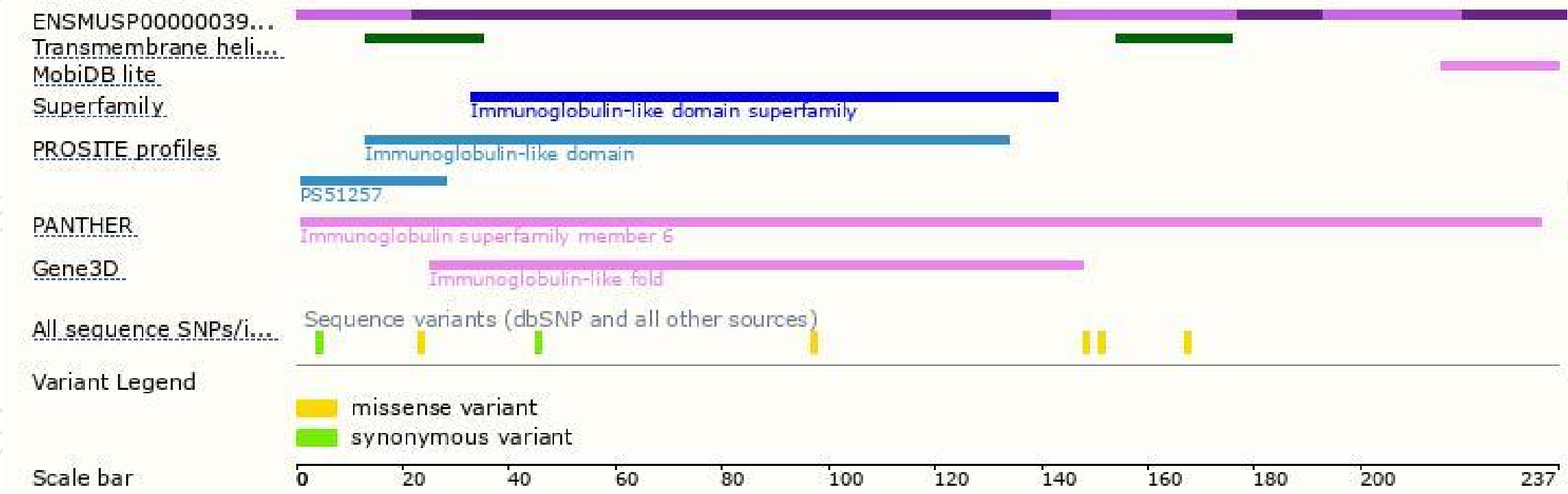
The strategy is based on the design of *Igsf6-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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