

***Fam83b* Cas9-KO Strategy**

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Project Overview

Project Name

Fam83b

Project type

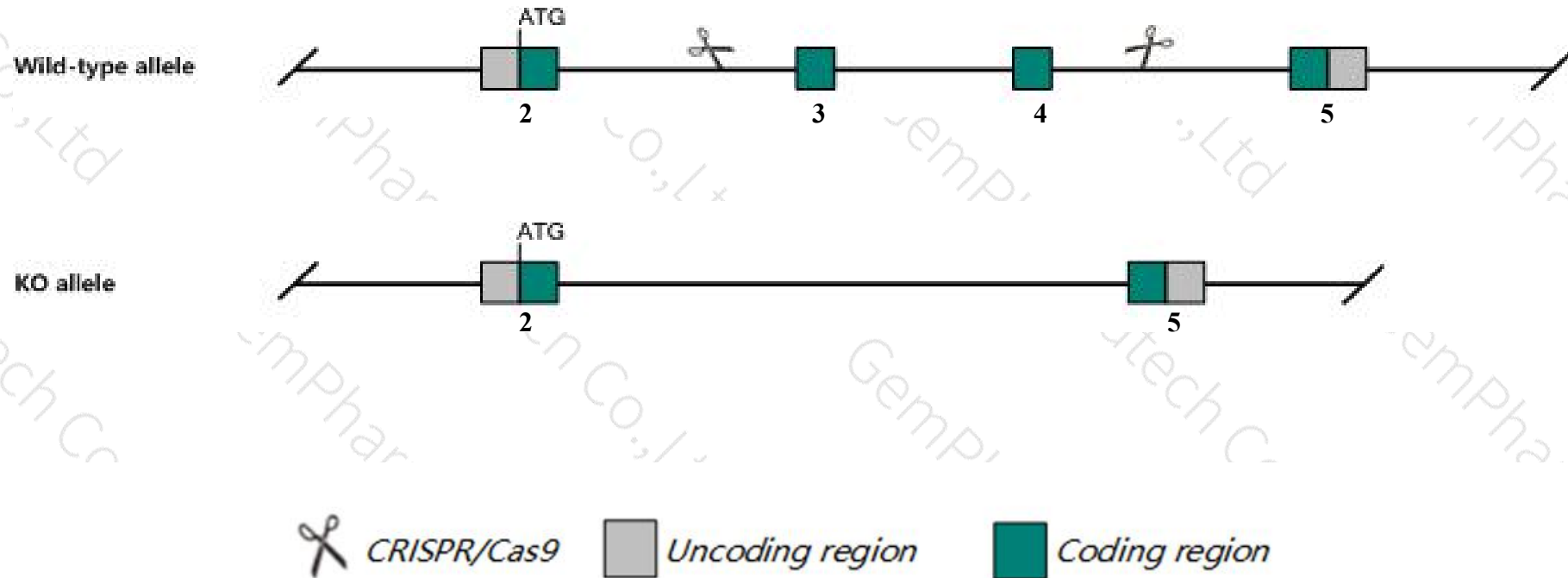
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fam83b* gene. The schematic diagram is as follows:



- The *Fam83b* gene has 2 transcripts. According to the structure of *Fam83b* gene, exon3-exon4 of *Fam83b*-202 (ENSMUST00000183437.7) transcript is recommended as the knockout region. The region contains 290bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fam83b* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Fam83b* gene is located on the Chr9. 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)

Fam83b family with sequence similarity 83, member B [*Mus musculus* (house mouse)]

Gene ID: 208994, updated on 14-Aug-2019

Summary

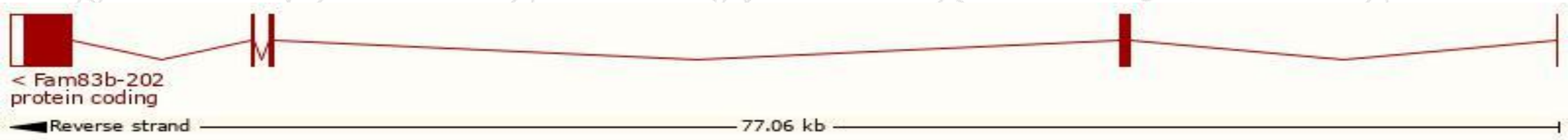
Official Symbol	Fam83b provided by MGI
Official Full Name	family with sequence similarity 83, member B provided by MGI
Primary source	MGI:MGI:2685362
See related	Ensembl:ENSMUSG00000032358
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Gm516; C530008M07Rik
Expression	Biased expression in large intestine adult (RPKM 1.4), small intestine adult (RPKM 1.1) and 8 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

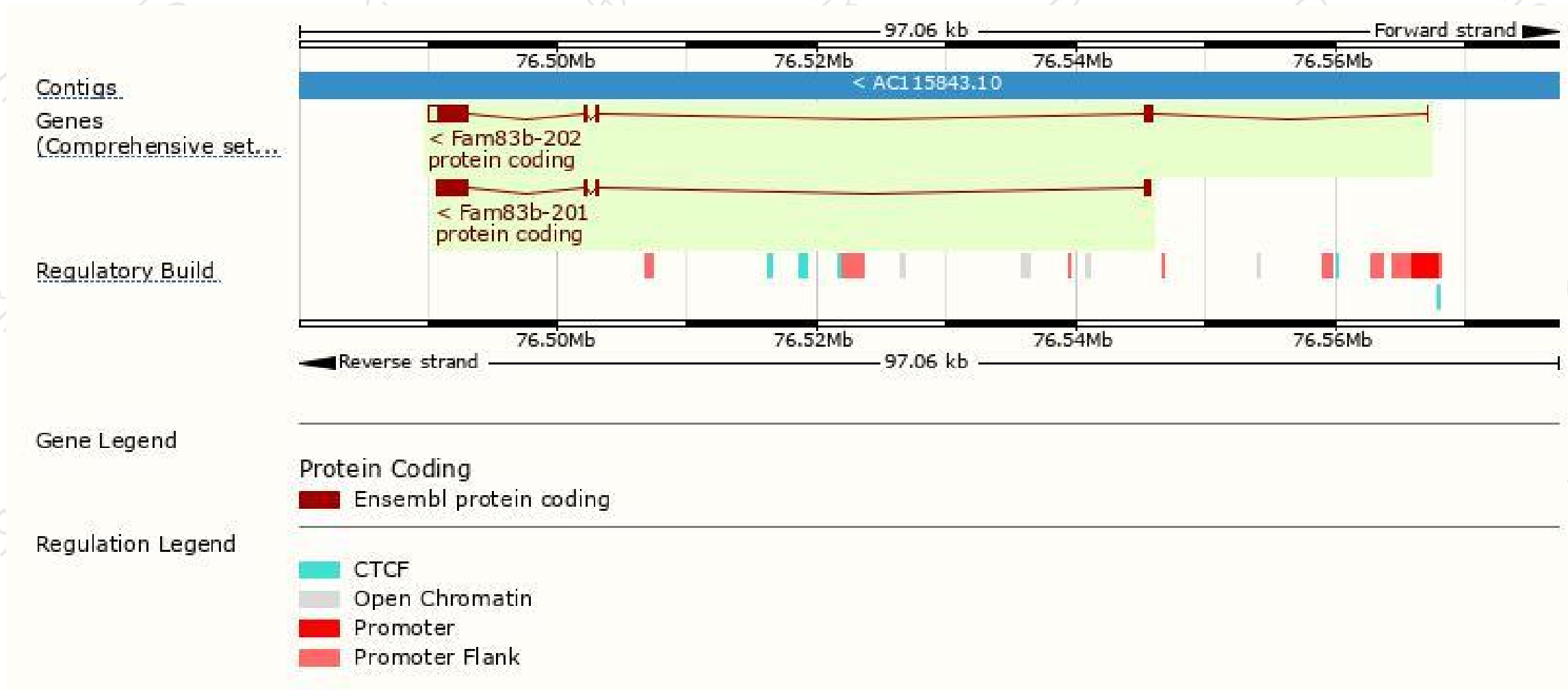
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fam83b-202	ENSMUST00000183437.7	3879	1012aa	Protein coding	CCDS40695	Q0VBM2	TSL:1 GENCODE basic APPRIS P1
Fam83b-201	ENSMUST00000098546.3	3136	1012aa	Protein coding	CCDS40695	Q0VBM2	TSL:1 GENCODE basic APPRIS P1

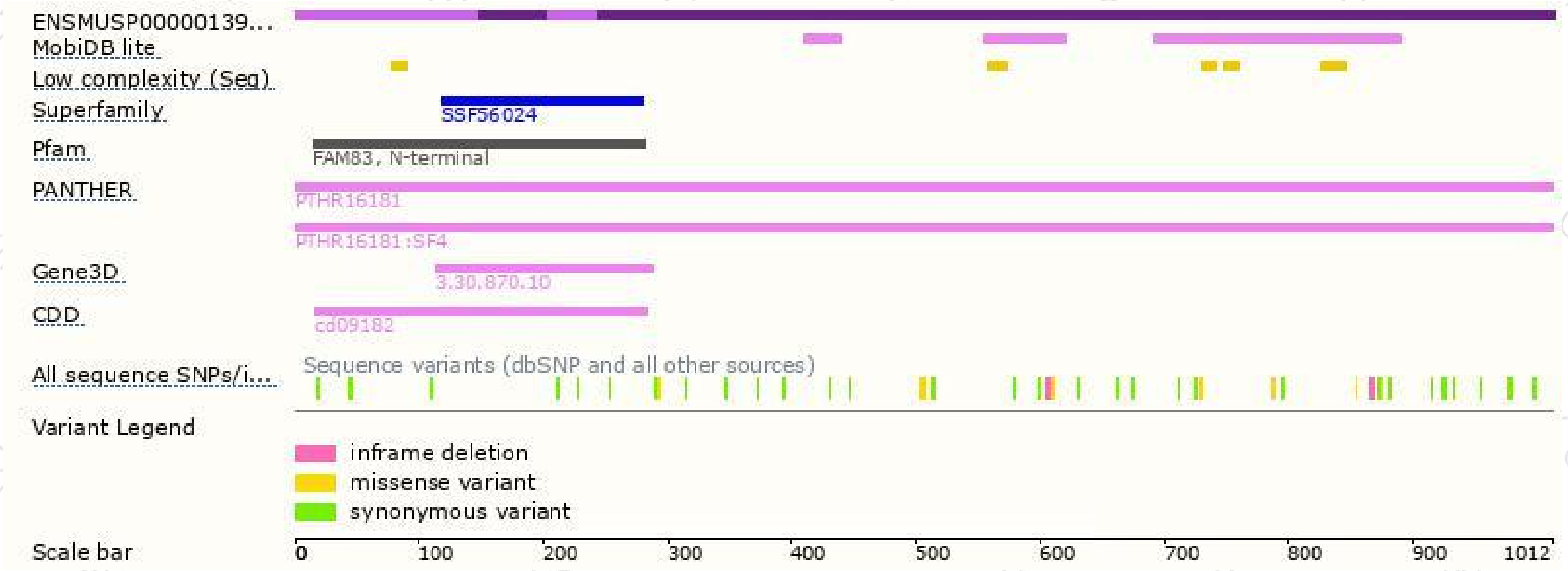
The strategy is based on the design of *Fam83b-202* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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