

Fbln7 Cas9-KO Strategy

Designer:

Jinling Wang

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2019-8-8

Project Overview

Project Name

Fbln7

Project type

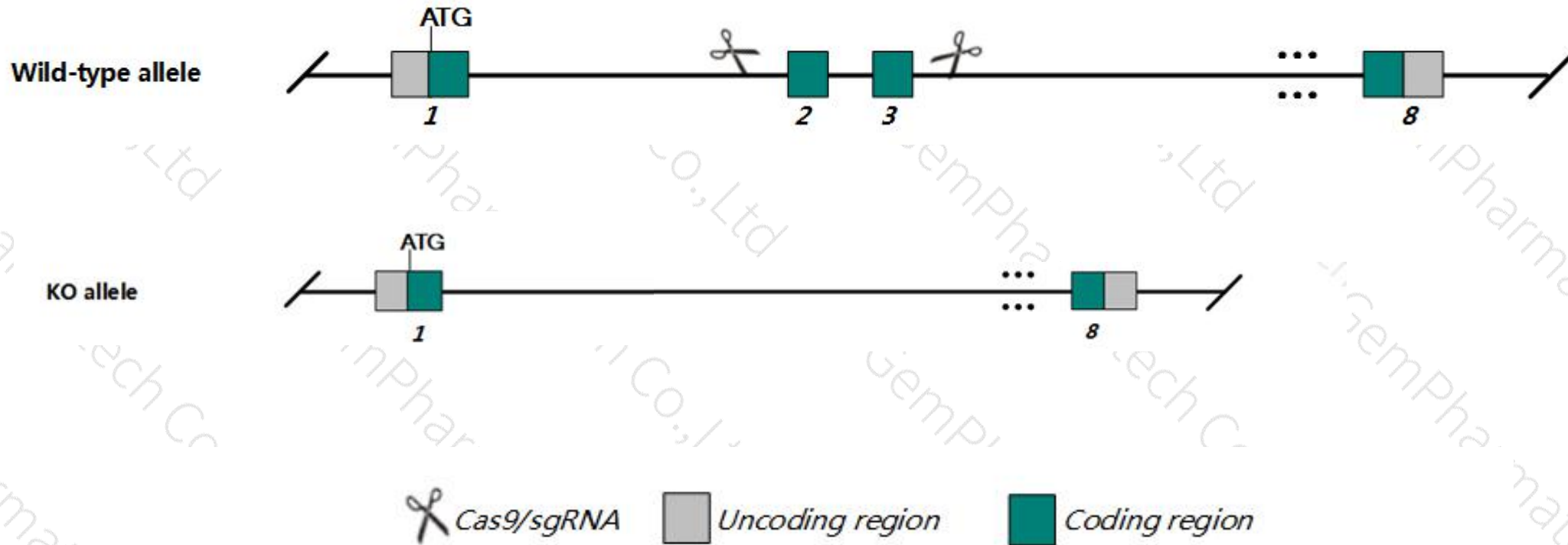
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fbln7* gene. The schematic diagram is as follows:



- The *Fbln7* gene has 2 transcripts. According to the structure of *Fbln7* gene, exon2-3 of *Fbln7-201* (ENSMUST00000028864.2) transcript is recommended as the knockout region. The region contains 331bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fbln7* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Fbln7* gene is located on the Chr2. 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)

Fbln7 fibulin 7 [Mus musculus (house mouse)]

Gene ID: 70370, updated on 31-Jan-2019

Summary



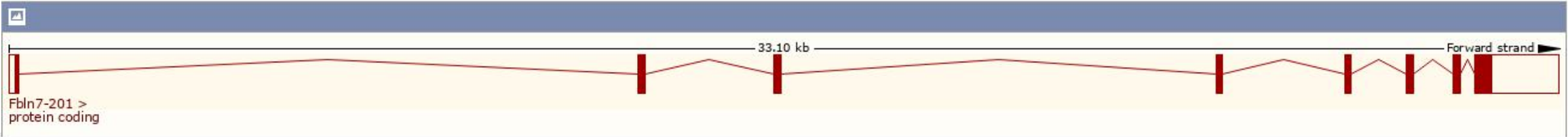
Official Symbol	Fbln7 provided by MGI
Official Full Name	fibulin 7 provided by MGI
Primary source	MGI:MGI:1917620
See related	Ensembl:ENSMUSG000000027386
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1600015H20Rik, AI464334, TM14
Expression	Broad expression in placenta adult (RPKM 6.7), kidney adult (RPKM 6.3) and 19 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

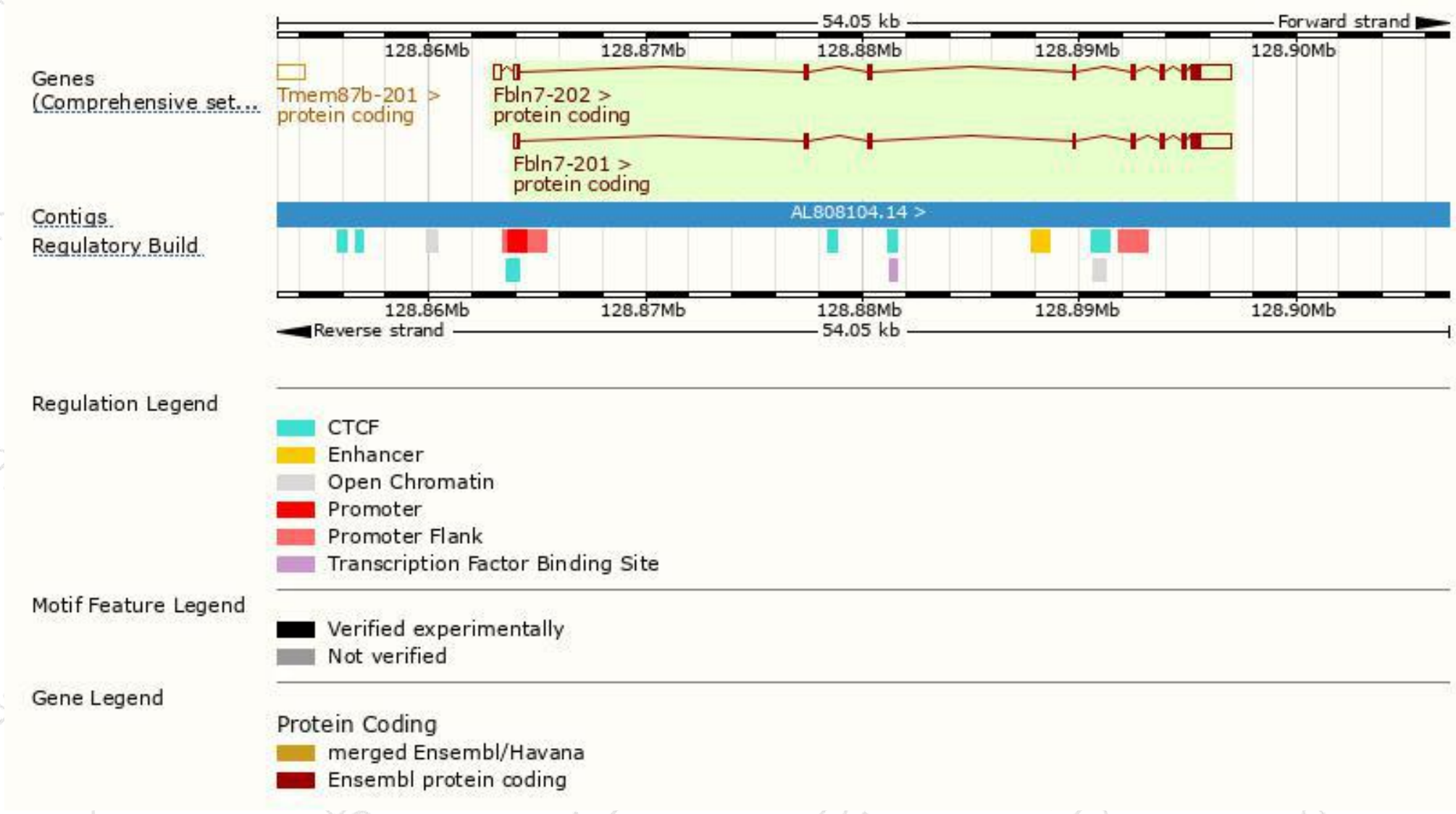
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fbln7-202	ENSMUST00000110324.7	3228	440aa	Protein coding	CCDS38235	Q501P1	TSL:5 GENCODE basic APPRIS P1
Fbln7-201	ENSMUST00000028864.2	2905	440aa	Protein coding	CCDS38235	Q501P1	TSL:1 GENCODE basic APPRIS P1

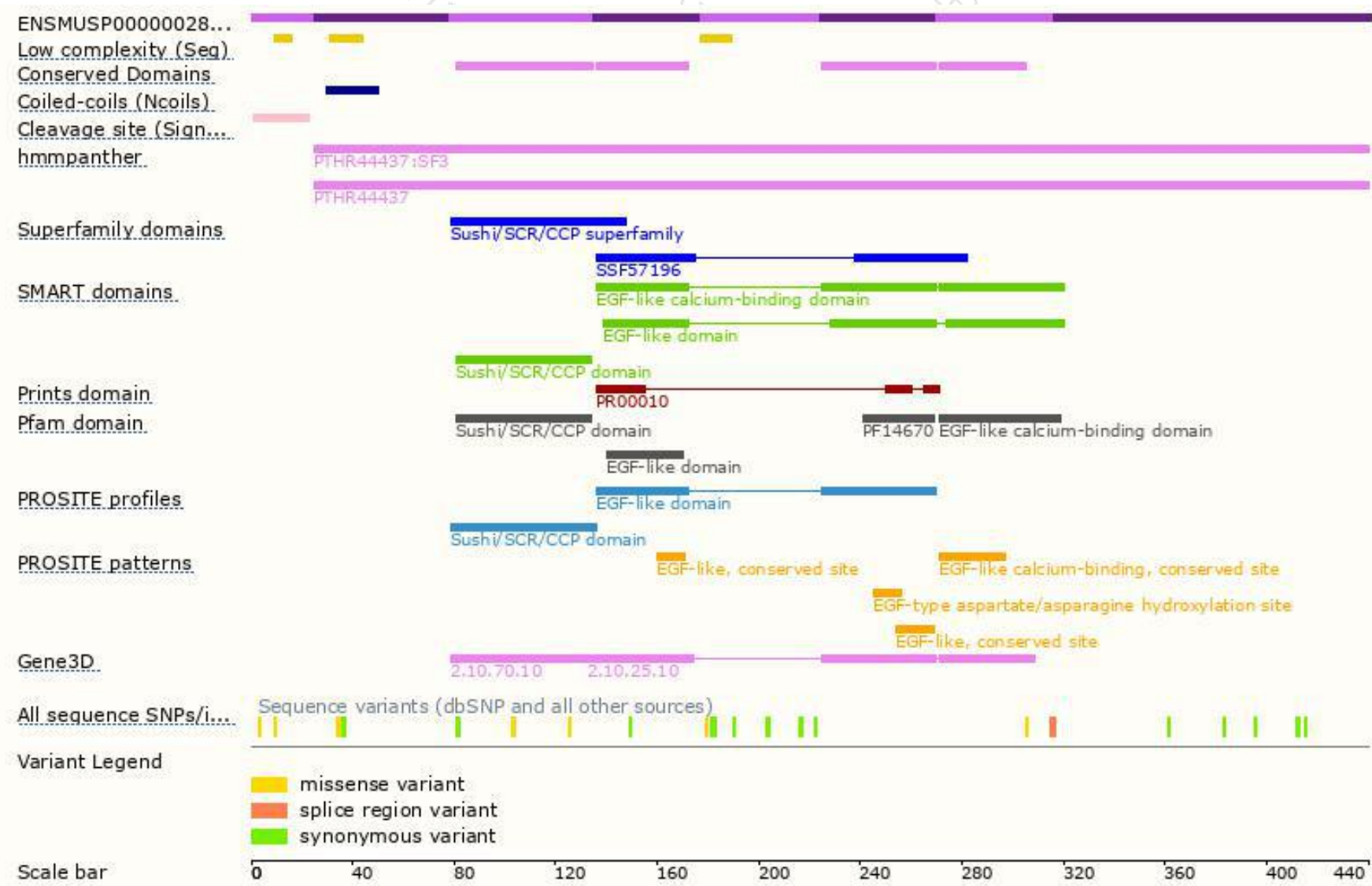
The strategy is based on the design of *Fbln7-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

Tel: 400-9660890

