

Lrrc8b Cas9-CKO Strategy

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Project Overview

Project Name

Lrrc8b

Project type

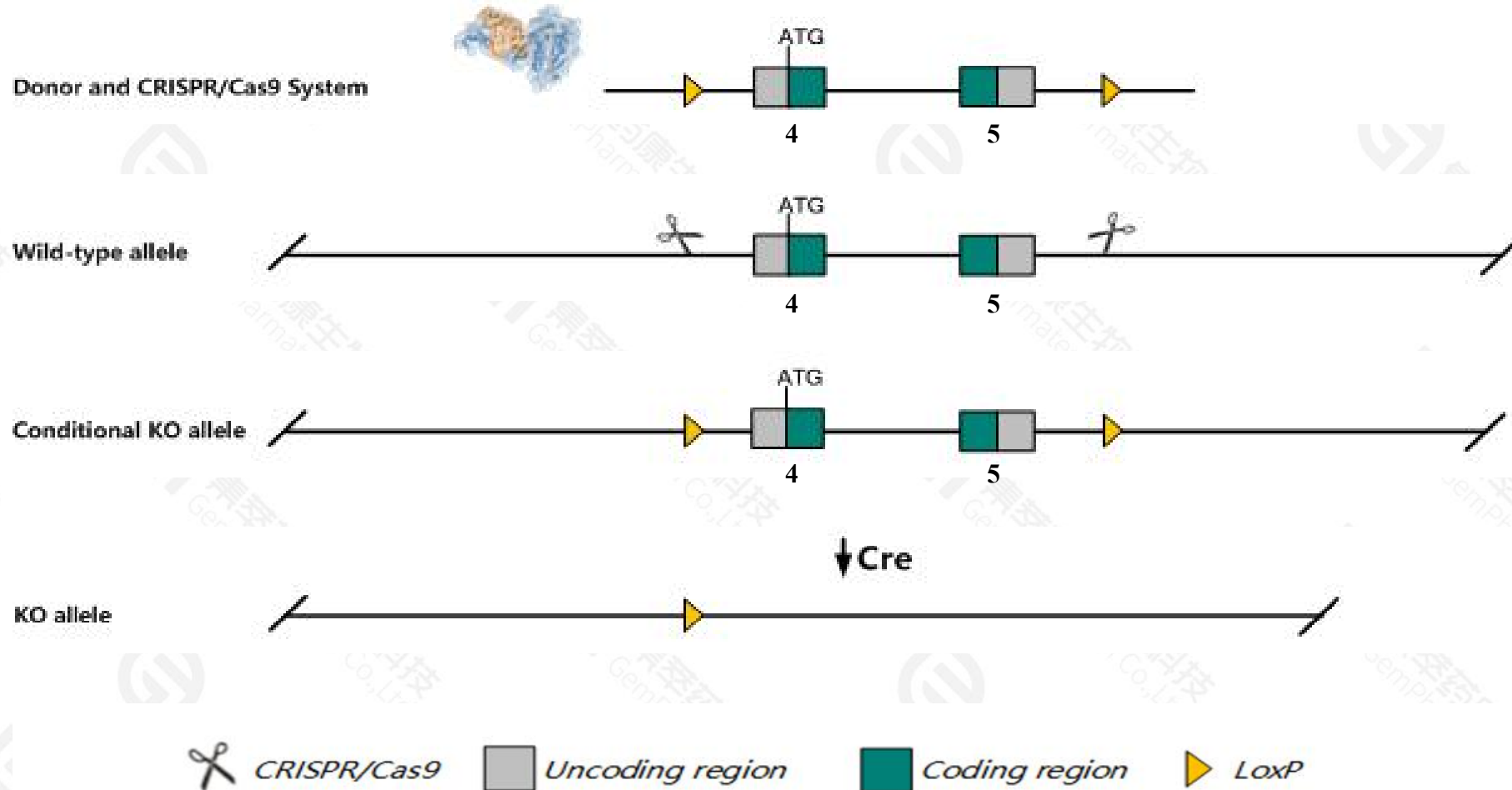
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lrrc8b* gene. The schematic diagram is as follows:



Technical routes

- The *Lrrc8b* gene has 1 transcript. According to the structure of *Lrrc8b* gene, exon4-exon5 of *Lrrc8b*-201(ENSMUST00000112707.3) transcript is recommended as the knockout region. The region contains all 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 *Lrrc8b* gene. The brief process is as follows: CRISPR/Cas9 system and Donor 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 flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

- The *Lrrc8b* gene is located on the Chr5. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Lrrc8b leucine rich repeat containing 8 family, member B [Mus musculus (house mouse)]

Gene ID: 433926, updated on 3-Jan-2021

Summary



Official Symbol Lrrc8b provided by [MGI](#)

Official Full Name leucine rich repeat containing 8 family, member B provided by [MGI](#)

Primary source [MGI:MGI:2141353](#)

See related [Ensembl:ENSMUSG00000070639](#)

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 2210408K08Rik, R7558, R75581, Ta-lrrp, mKIAA0231

Expression Biased expression in testis adult (RPKM 142.7), adrenal adult (RPKM 13.7) and 5 other tissues [See more](#)

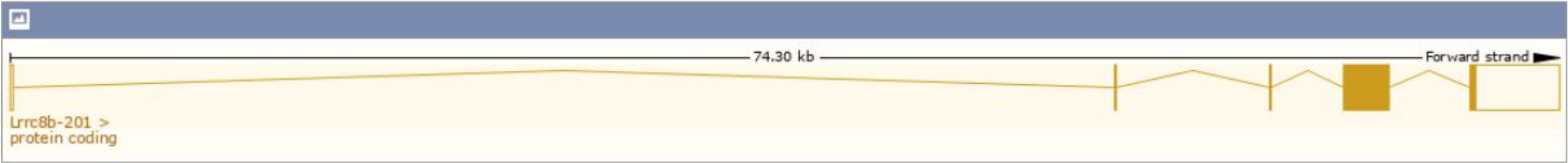
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

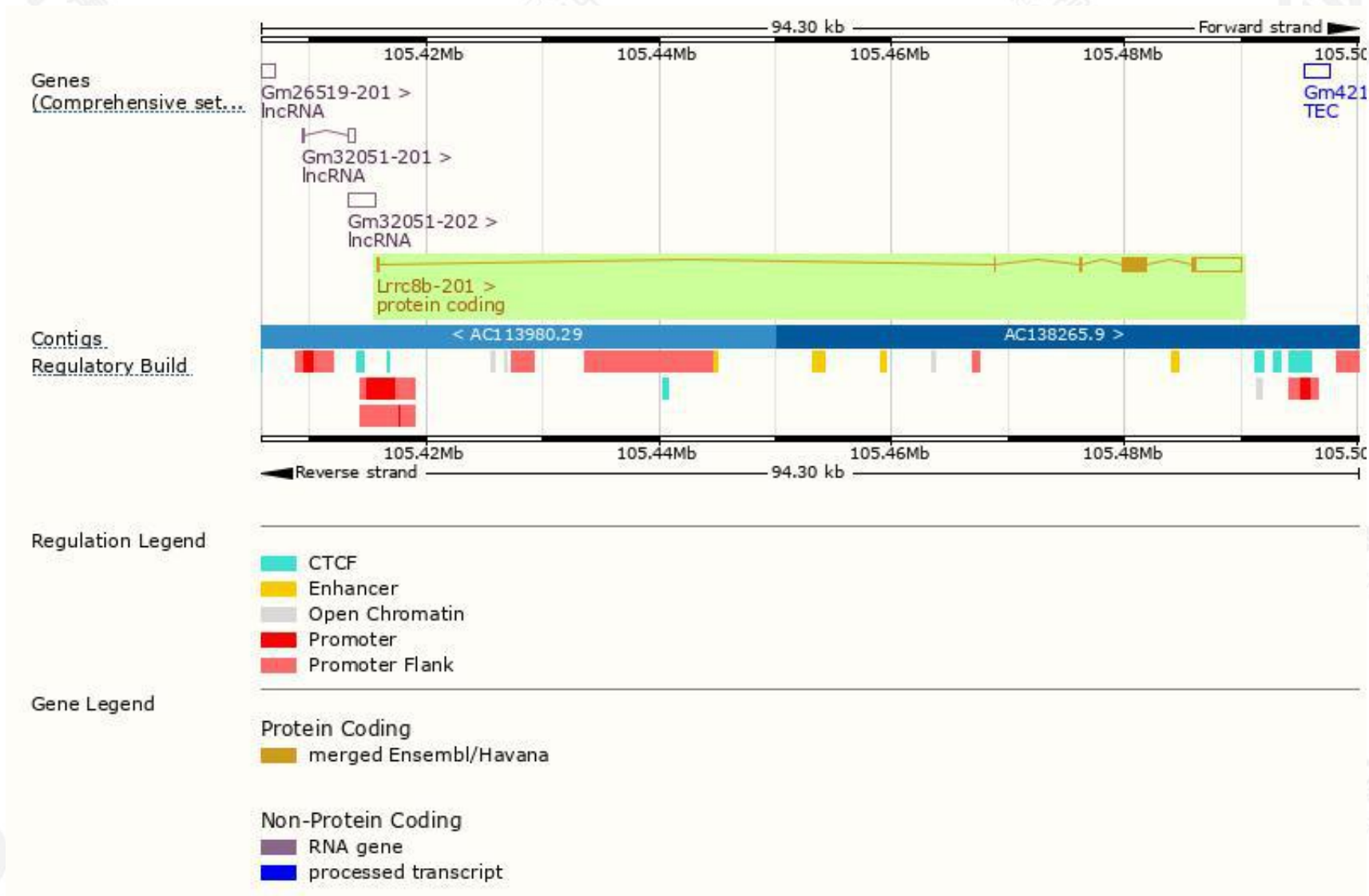
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lrrc8b-201	ENSMUST00000112707.3	6726	803aa	Protein coding	CCDS39193		TSL:1 , GENCODE basic , APPRIS P1 ,

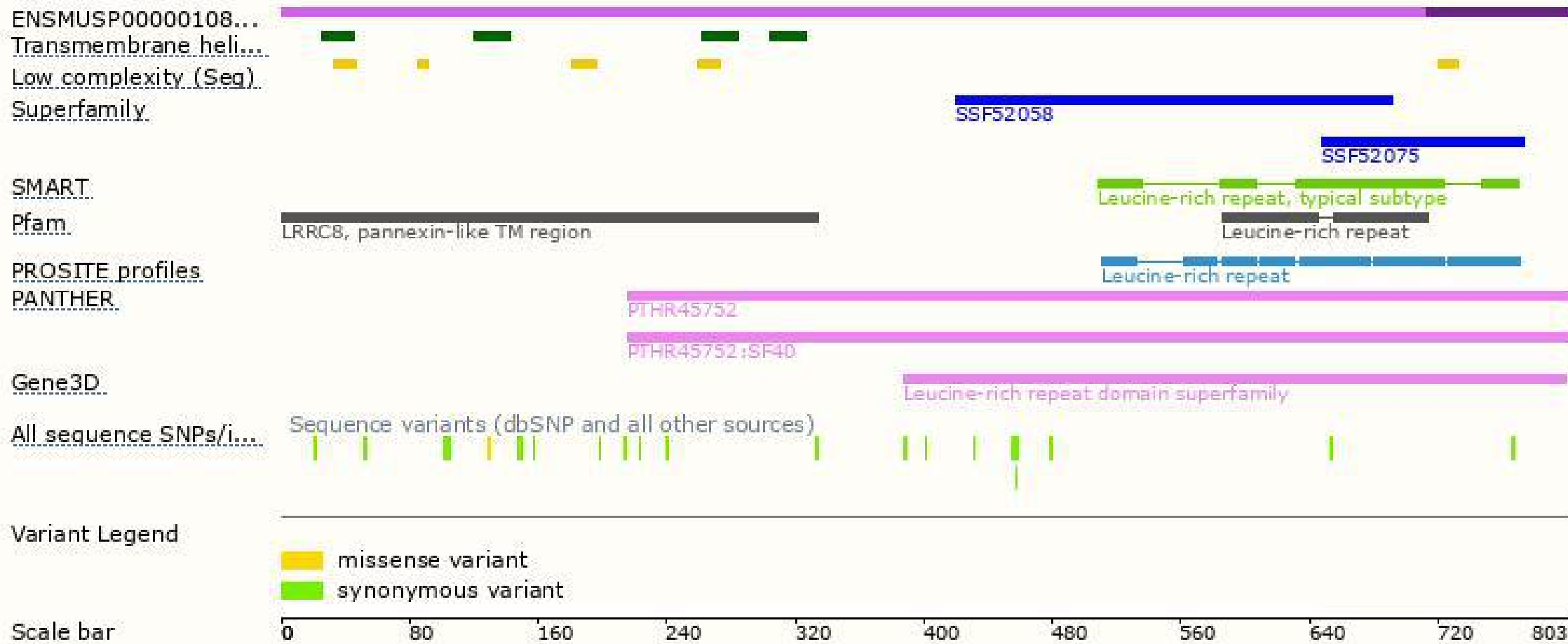
The strategy is based on the design of *Lrrc8b-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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