

Lrrn4 Cas9-KO Strategy

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Project Overview

Project Name

Lrrn4

Project type

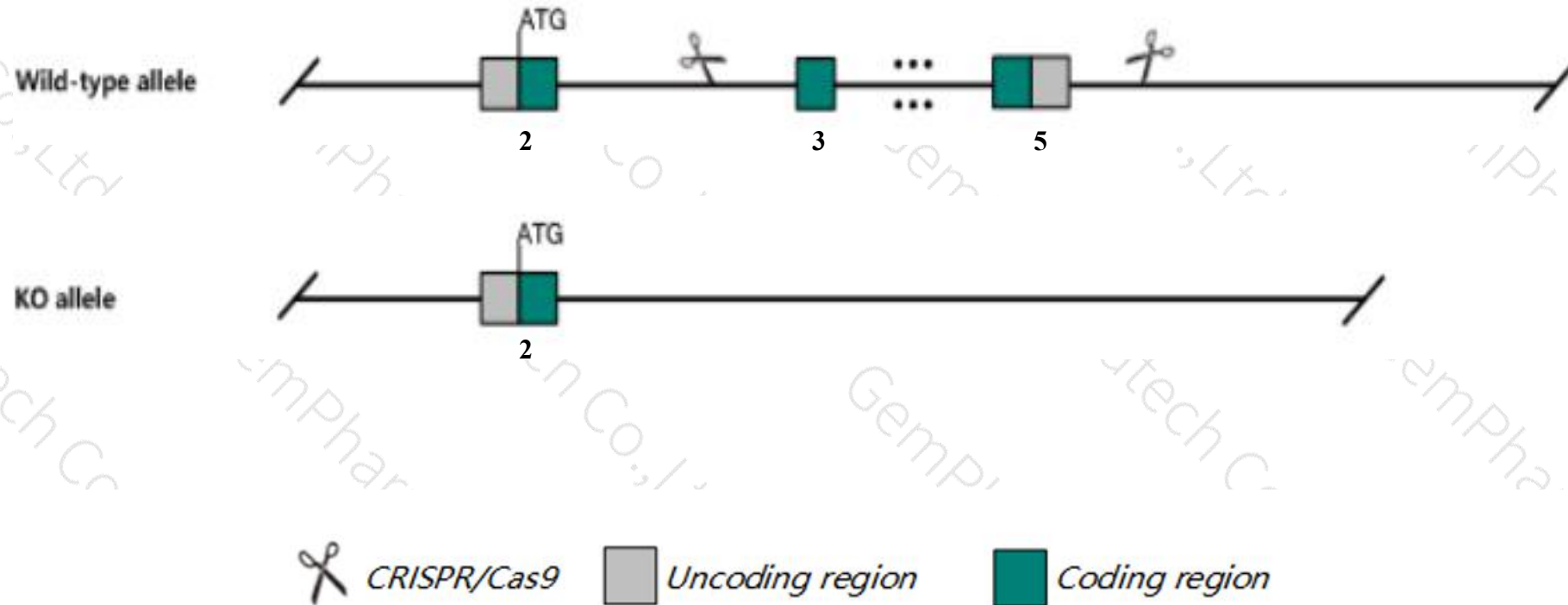
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lrrn4* gene. The schematic diagram is as follows:



- The *Lrrn4* gene has 1 transcript. According to the structure of *Lrrn4* gene, exon3-exon5 of *Lrrn4-201*(ENSMUST00000049787.2) transcript is recommended as the knockout region. The region contains 1541bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Lrrn4* 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.

- According to the existing MGI data, homozygous null mutant mice exhibit impaired memory retention in hippocampus-dependent learning tasks such as contextual conditioning and spatial learning.
- The *Lrrn4* 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)

Lrrn4 leucine rich repeat neuronal 4 [Mus musculus (house mouse)]

Gene ID: 320974, updated on 13-Mar-2020

Summary



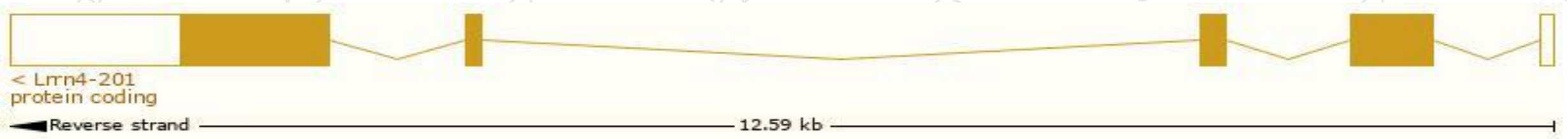
Official Symbol	Lrrn4 provided by MGI
Official Full Name	leucine rich repeat neuronal 4 provided by MGI
Primary source	MGI:MGI:2445154
See related	Ensembl:ENSMUSG00000043110
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	B430119L13Rik, Nlrr4
Expression	Broad expression in lung adult (RPKM 11.4), ovary adult (RPKM 9.1) and 15 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

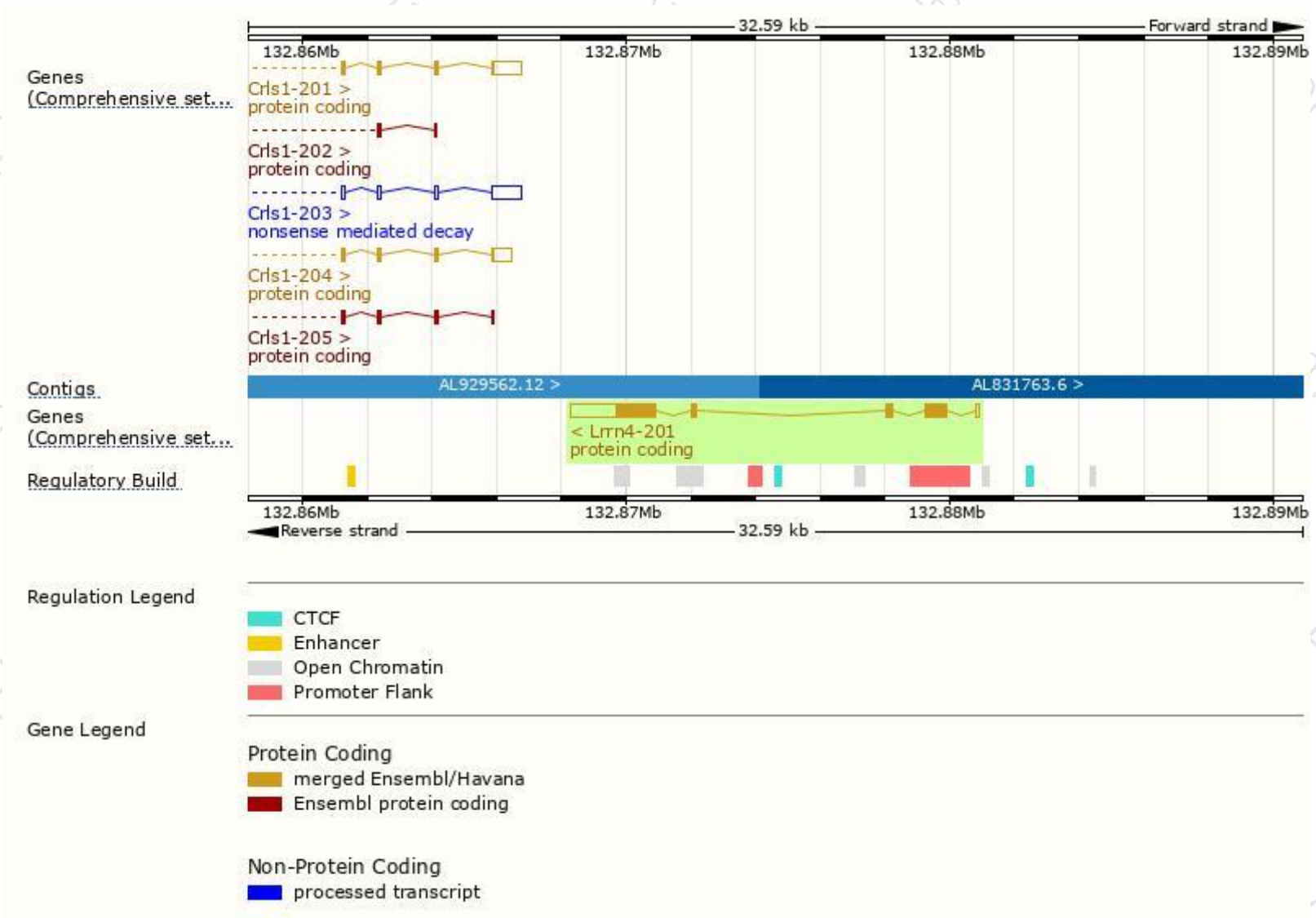
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lrrn4-201	ENSMUST00000049787.2	3715	733aa	Protein coding	CCDS16780	P59383	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Lrrn4-201* transcript, the transcription is shown below:



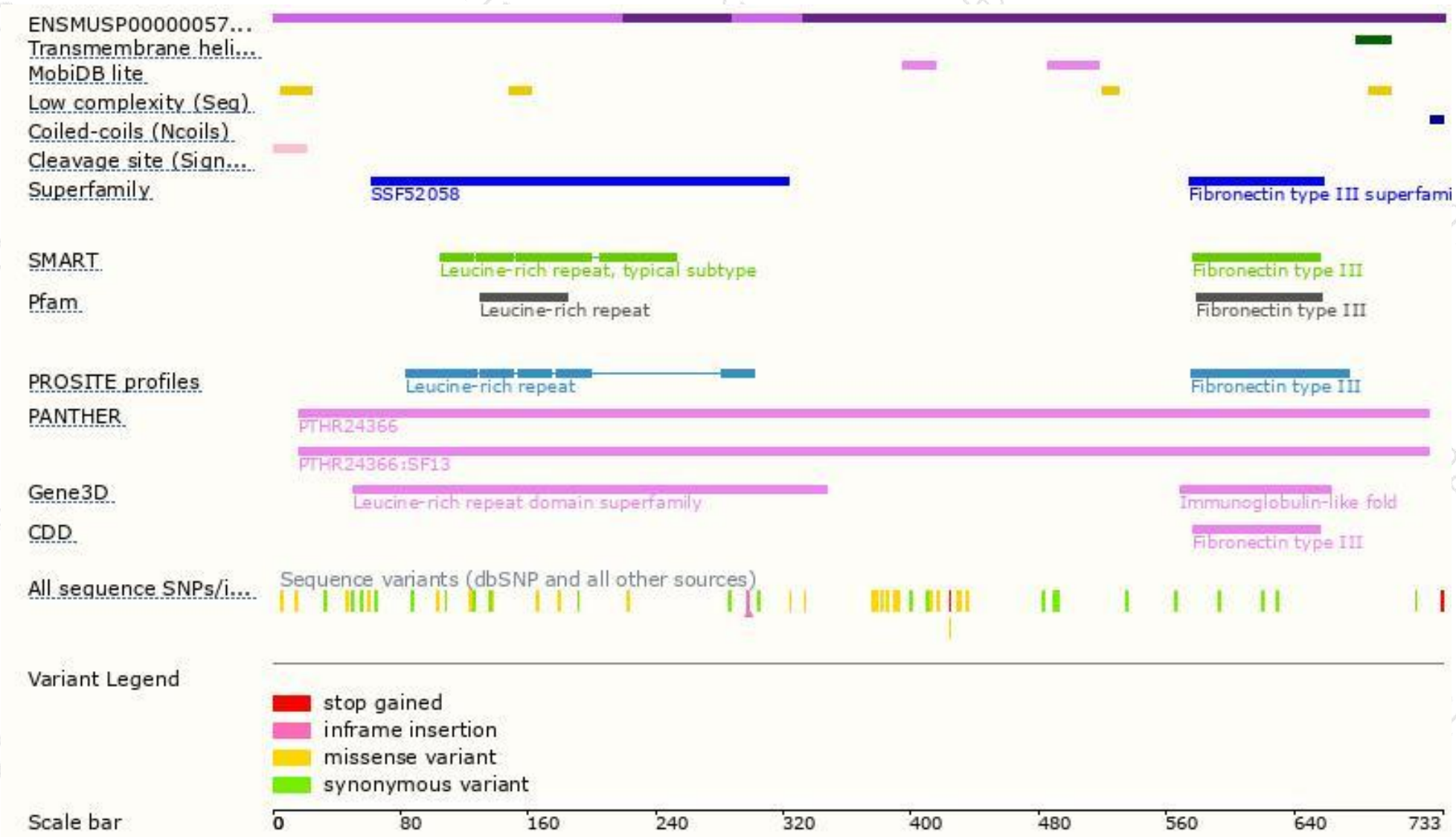
Genomic location distribution



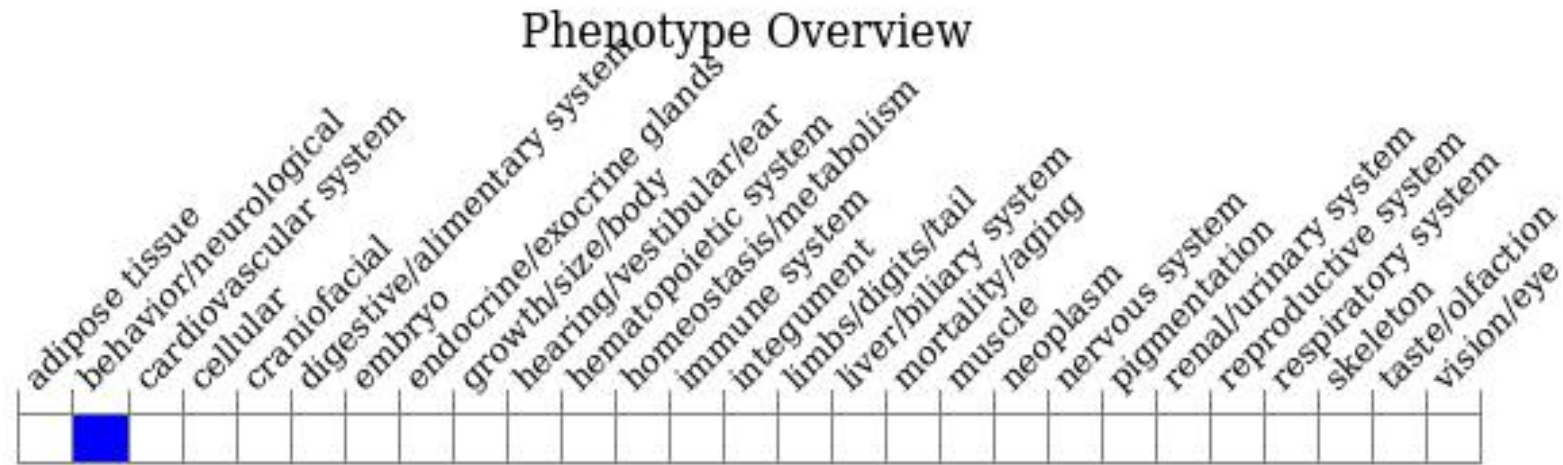
Protein domain



集萃药康
GemPharmatech



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

According to the existing MGI data, homozygous null mutant mice exhibit impaired memory retention in hippocampus- dependent learning tasks such as contextual conditioning and spatial learning.

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

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