

Lym4 Cas9-KO Strategy

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Project Overview

Project Name

Lym4

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lym4* gene. The schematic diagram is as follows:



- The *Lym4* gene has 1 transcript. According to the structure of *Lym4* gene, exon2 of *Lym4-201* (ENSMUST00000053265.7) transcript is recommended as the knockout region. The region contains 121bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Lym4* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Lym4* gene is located on the Chr13. 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.
- The partial intron of *Gm48765* gene will be deleted together in this strategy.
- 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)

Lyrm4 LYR motif containing 4 [Mus musculus (house mouse)]

Gene ID: 380840, updated on 13-Mar-2020

Summary



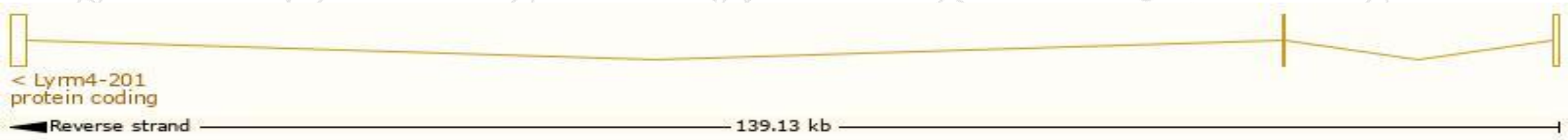
Official Symbol	Lyrm4 provided by MGI
Official Full Name	LYR motif containing 4 provided by MGI
Primary source	MGI:MGI:2683538
See related	Ensembl:ENSMUSG00000046573
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	BC034664, Gm903
Expression	Ubiquitous expression in CNS E18 (RPKM 6.6), cortex adult (RPKM 6.3) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

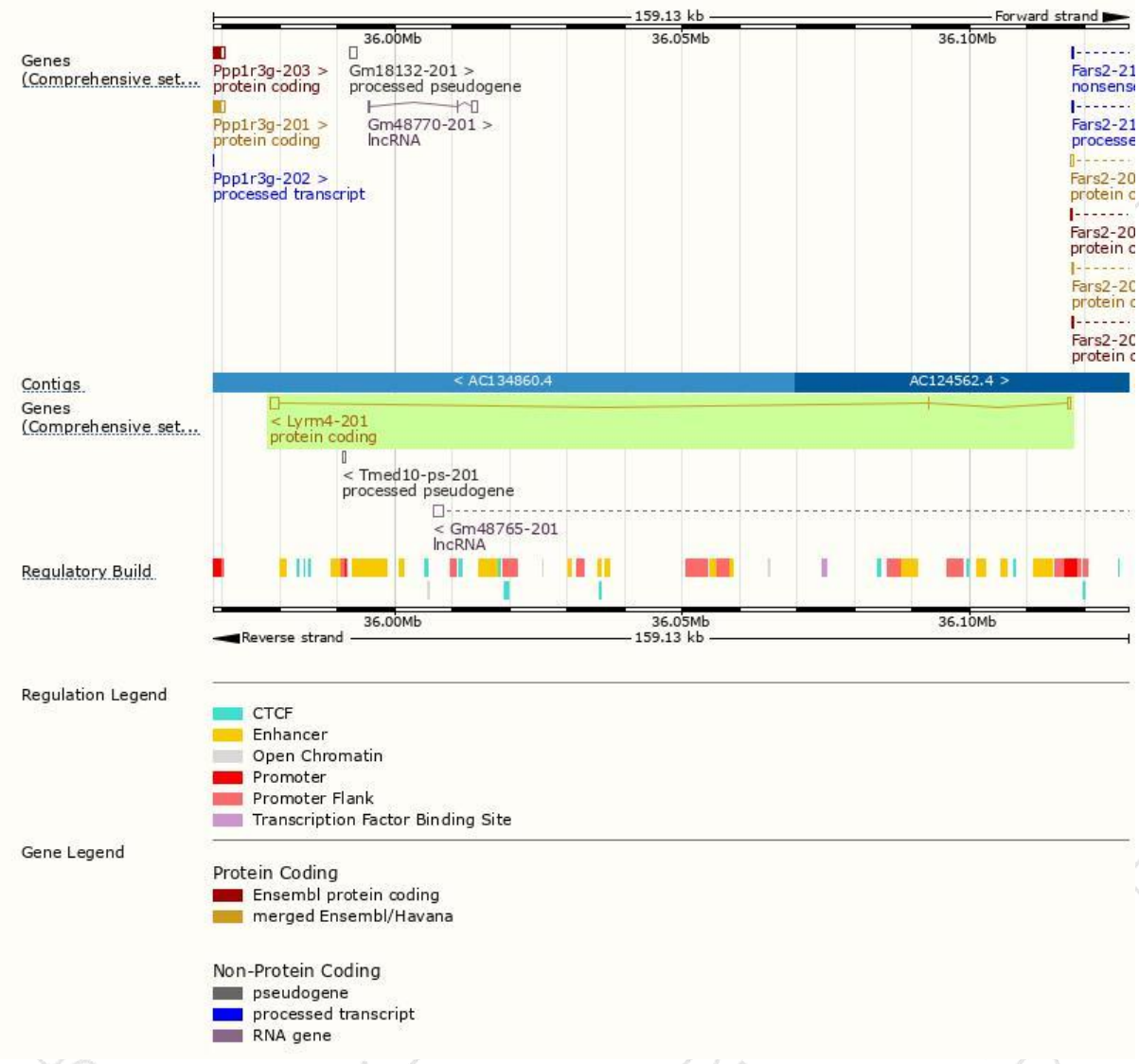
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lyrm4-201	ENSMUST00000053265.7	2086	91aa	Protein coding	CCDS36633	Q8K215	TSL:1 GENCODE basic APPRIS P1

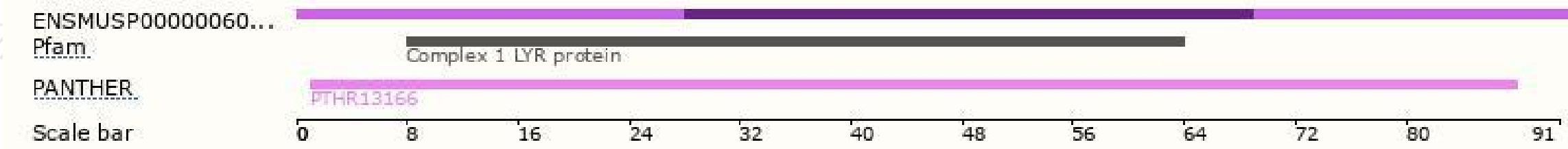
The strategy is based on the design of *Lyrm4-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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