

Lcn10 Cas9-KO Strategy

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Project Overview

Project Name

Lcn10

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lcn10* gene. The schematic diagram is as follows:



- The *Lcn10* gene has 1 transcript. According to the structure of *Lcn10* gene, exon2-exon4 of *Lcn10-201* (ENSMUST00000058912.2) transcript is recommended as the knockout region. The region contains 319bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Lcn10* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Lcn10* 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)

Lcn10 lipocalin 10 [Mus musculus (house mouse)]

Gene ID: 332578, updated on 31-Jan-2019

Summary



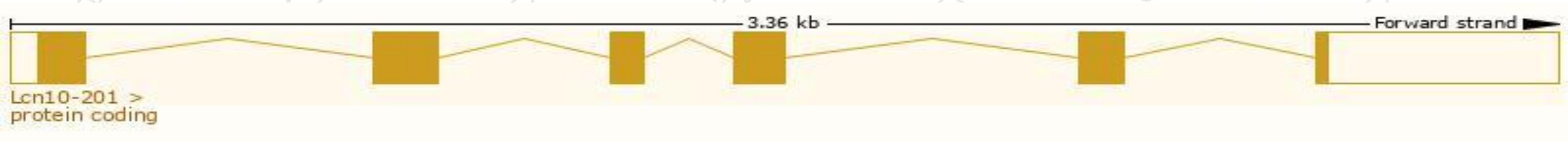
Official Symbol	Lcn10 provided by MGI
Official Full Name	lipocalin 10 provided by MGI
Primary source	MGI:MGI:1925000
See related	Ensembl:ENSMUSG00000047356
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	9230112J07Rik
Expression	Restricted expression toward genital fat pad adult (RPKM 356.4) See more
Orthologs	human all

Transcript information (Ensembl)

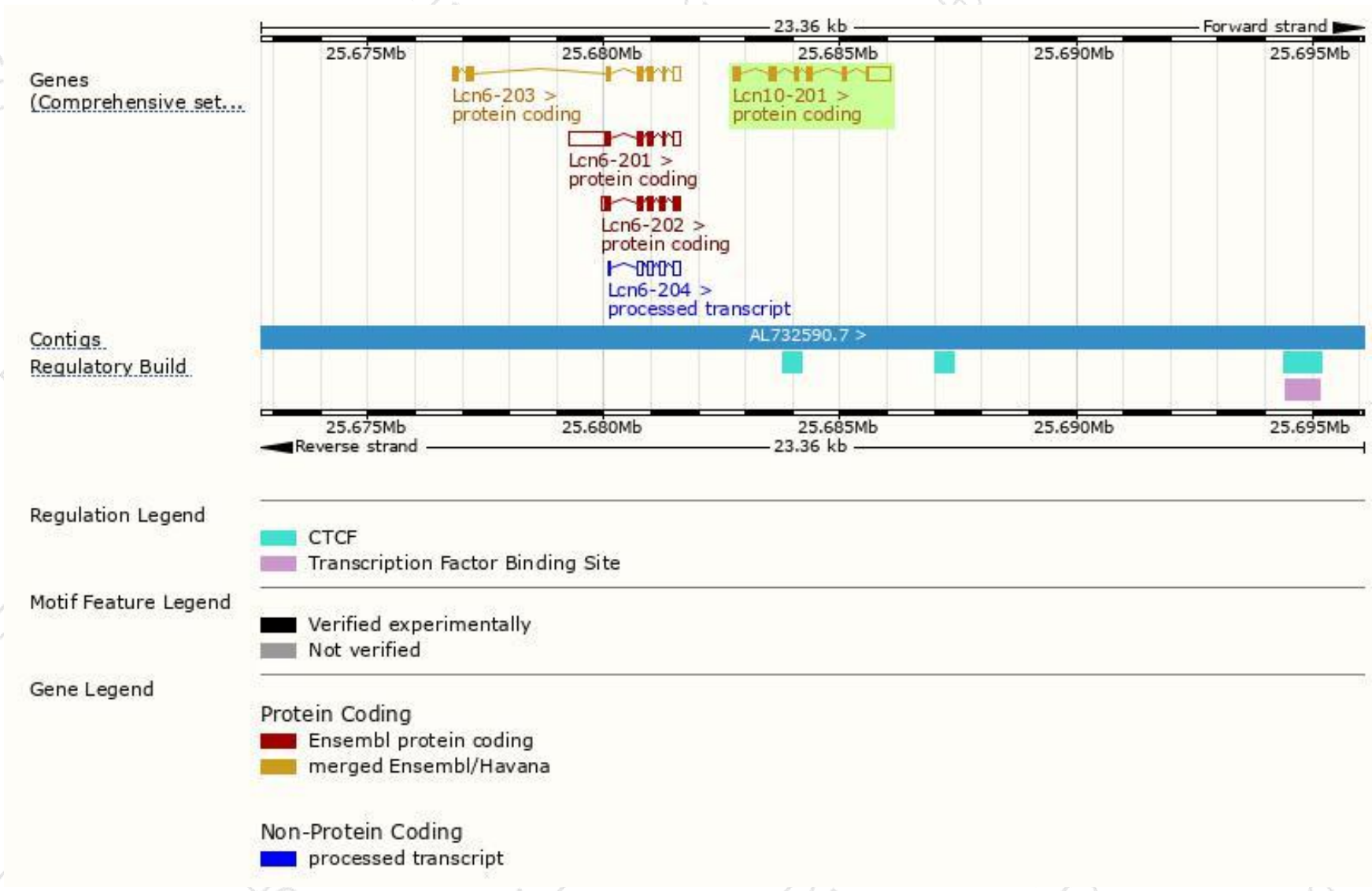
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Lcn10-201	ENSMUST00000058912.2	1106	182aa	Protein coding	CCDS15787	Q0VBF7 Q810Z1	TSL:1 GENCODE basic APPRIS P1

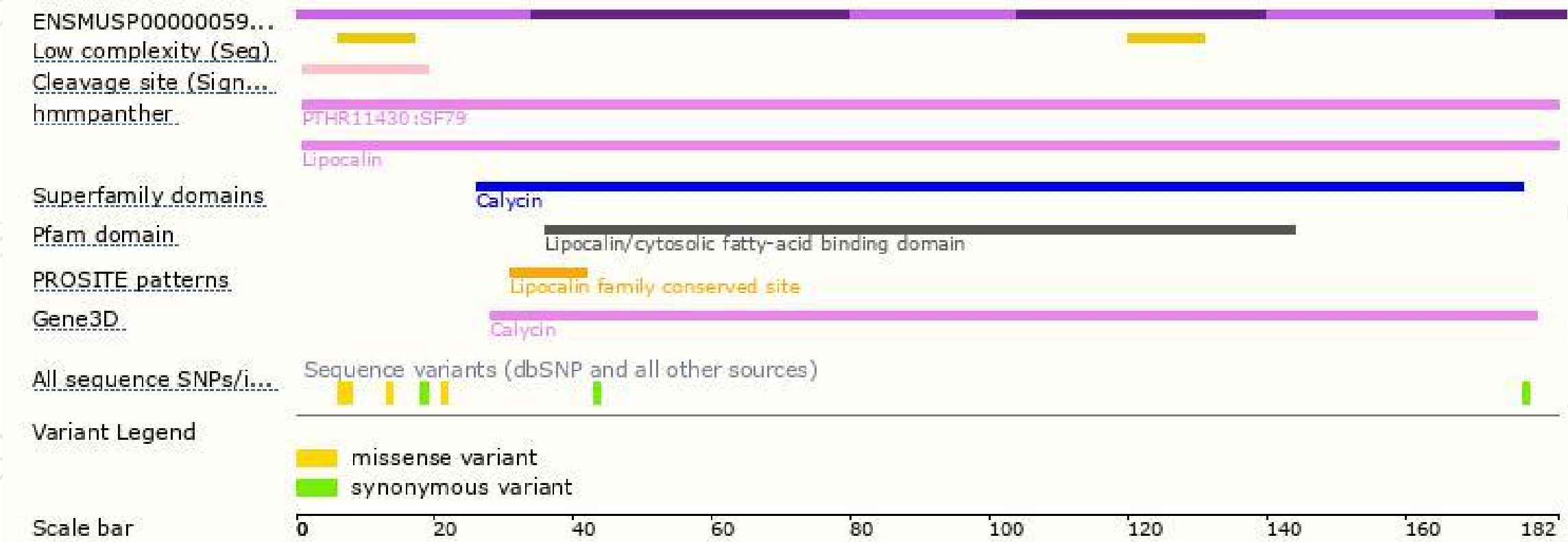
The strategy is based on the design of *Lcn10-201* transcript, The transcription is shown below



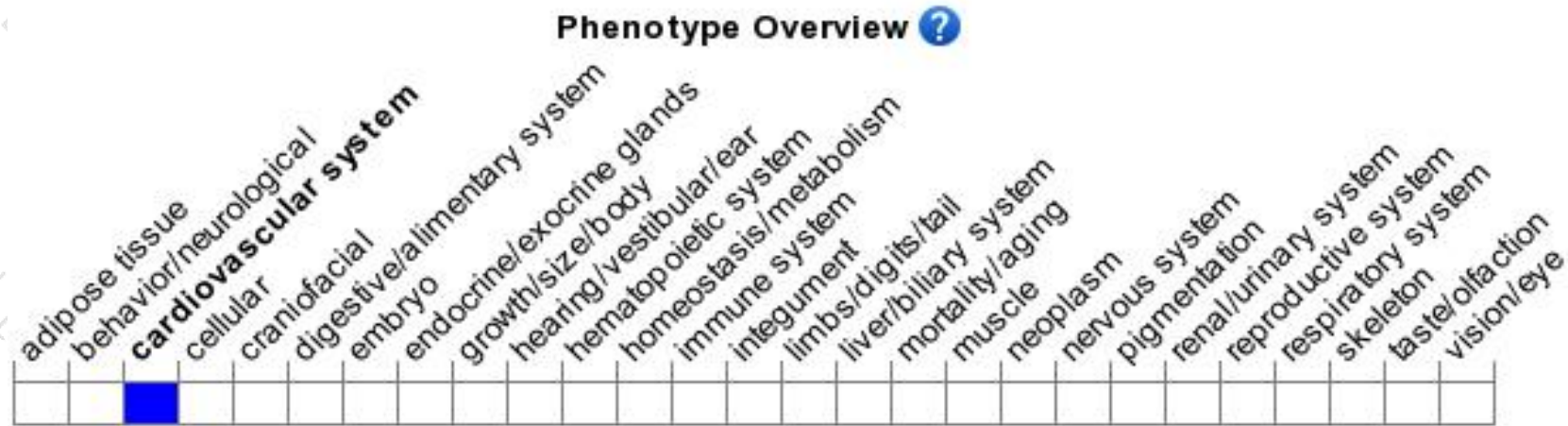
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



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

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

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