

Ch25h Cas9-KO Strategy

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Project Overview

Project Name

Ch25h

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ch25h* gene. The schematic diagram is as follows:



- The *Ch25h* gene has 1 transcript. According to the structure of *Ch25h* gene, exon1 of *Ch25h-201* (ENSMUST00000050562.5) 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 *Ch25h* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit increased IgG2a and IgA in the sera, lungs, and intestinal mucosa and increased IgG2b and IgG3 in the intestinal mucosa.
- The KO region contains functional region of the *Gm26902* gene. Knockout the region will affect the function of *Gm26902* gene.
- The *Ch25h* gene is located on the Chr19. 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)

Ch25h cholesterol 25-hydroxylase [Mus musculus (house mouse)]

Gene ID: 12642, updated on 19-Mar-2019

Summary



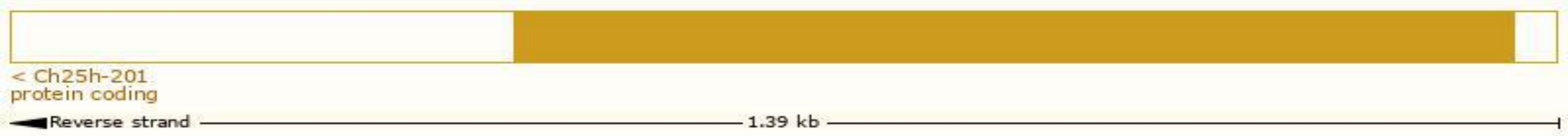
Official Symbol	Ch25h provided by MGI
Official Full Name	cholesterol 25-hydroxylase provided by MGI
Primary source	MGI:MGI:1333869
See related	Ensembl:ENSMUSG00000050370
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI462618, m25OH
Orthologs	human all

Transcript information (Ensembl)

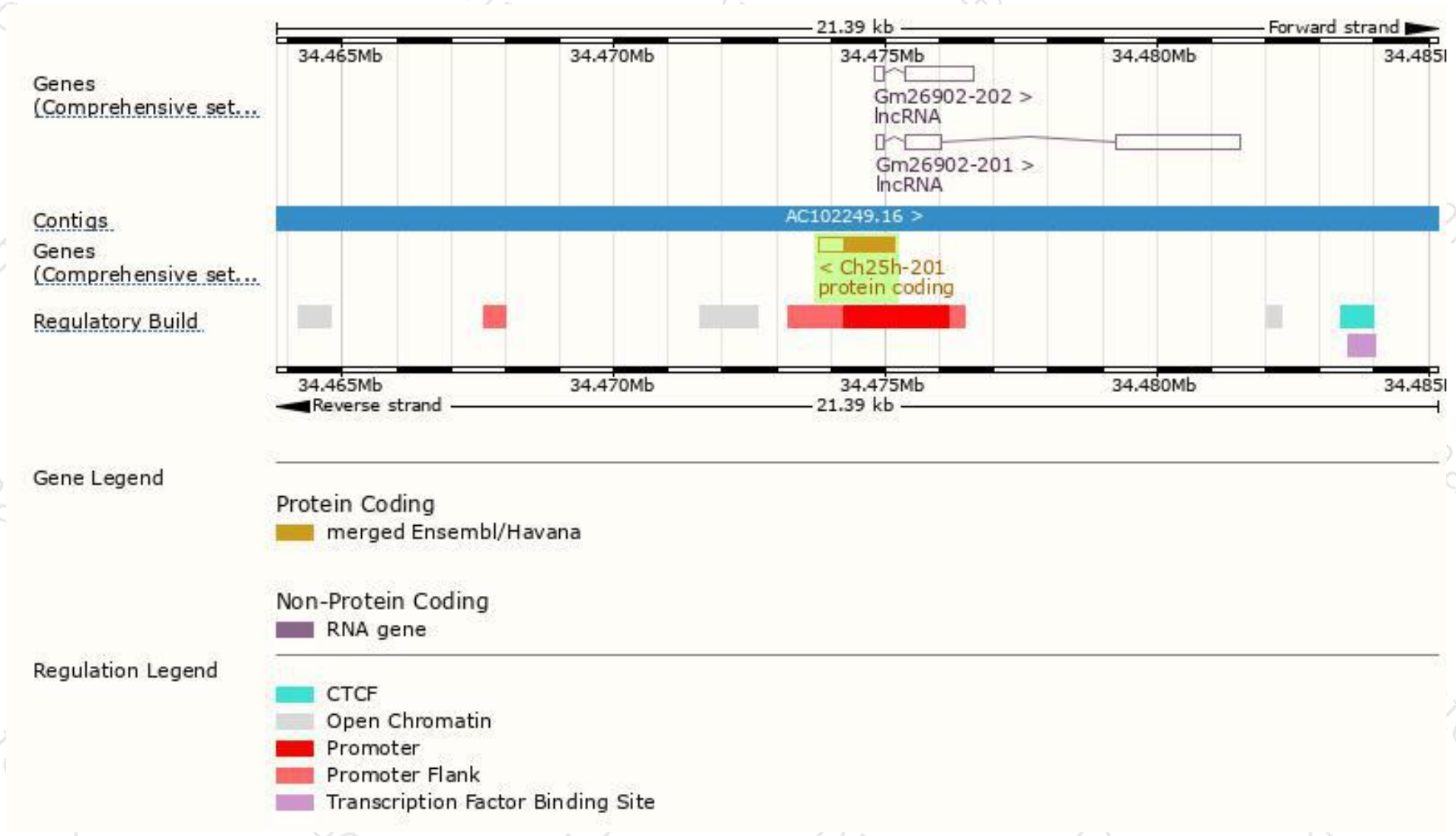
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ch25h-201	ENSMUST00000050562.5	1387	298aa	Protein coding	CCDS29759	Q9Z0F5	TSL:NA GENCODE basic APPRIS P1

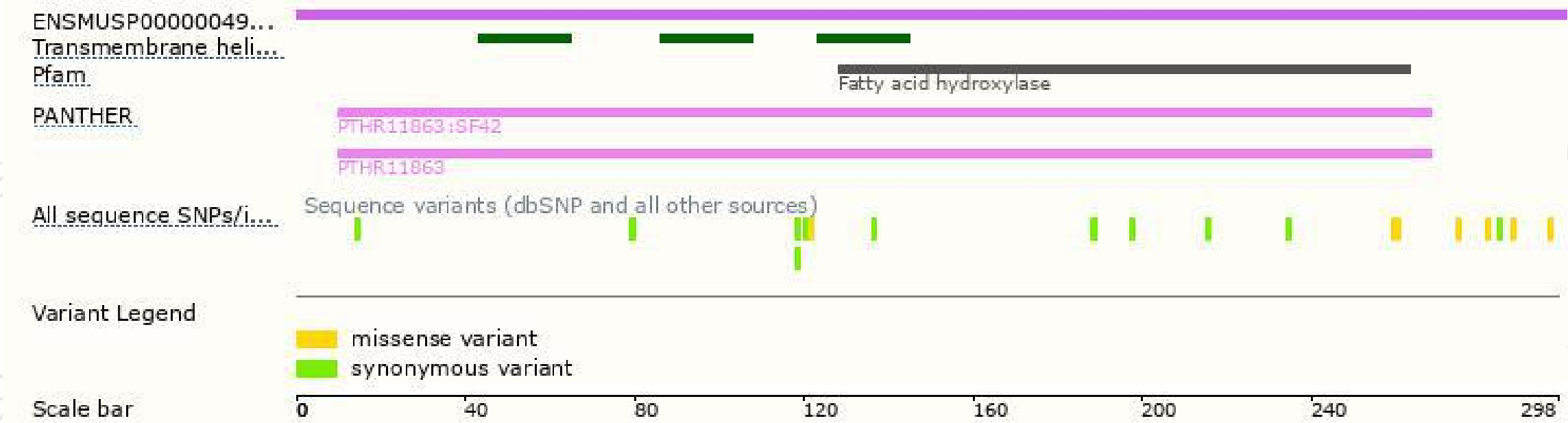
The strategy is based on the design of *Ch25h-201* transcript, The transcription is shown below



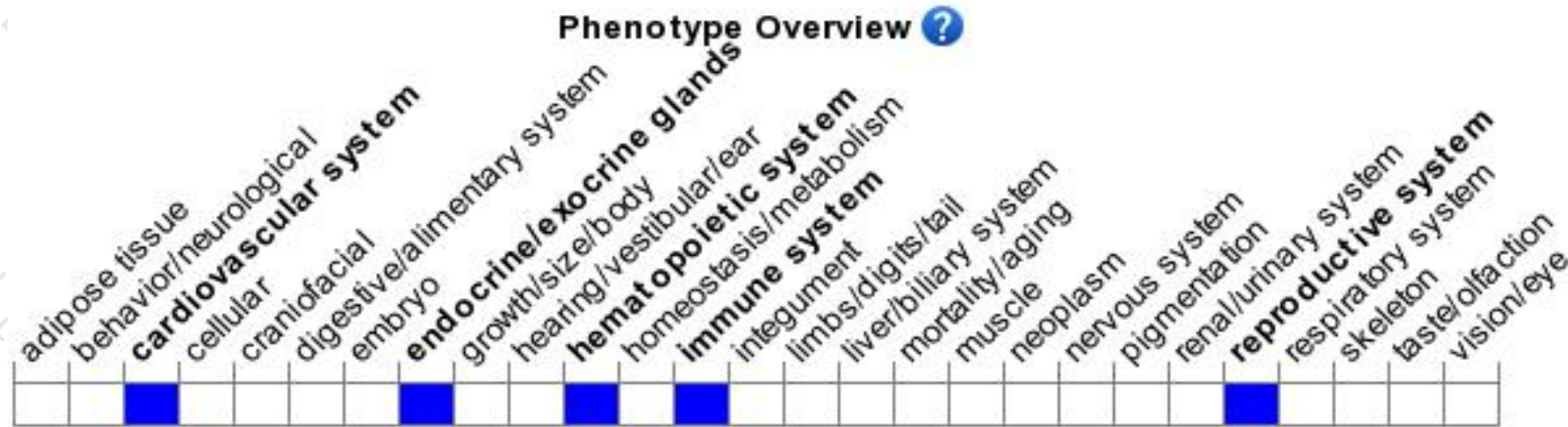
Genomic location distribution



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/>).

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit increased IgG2a and IgA in the sera, lungs, and intestinal mucosa and increased IgG2b and IgG3 in the intestinal mucosa.

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

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