Smyd2 Cas9-CKO Strategy

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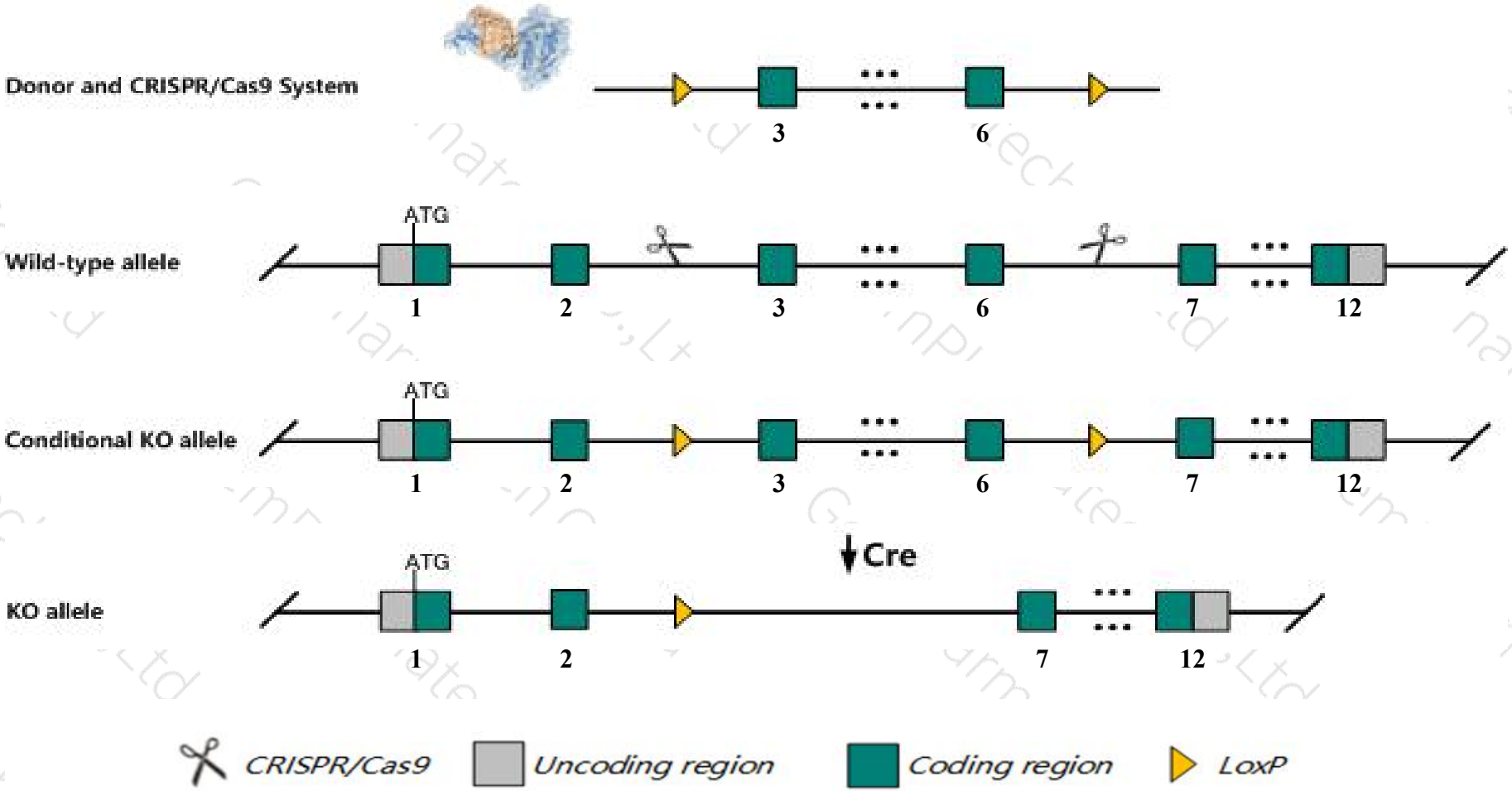
Project Overview

Project Name	***Smyd2***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Smyd2* gene. The schematic diagram is as follows:



Technical routes

- The *Smyd2* gene has 4 transcripts. According to the structure of *Smyd2* gene, exon3-exon6 of *Smyd2-201* (ENSMUST00000027897.7) transcript is recommended as the knockout region.The region contains 365bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Smyd2* 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.

Notice

- According to the existing MGI data,Mice homozygous for a targeted allele exhibit increased circulating total and LDL cholesterol levels and decreased circulating sodium and chloride levels.

- The *Smyd2* gene is located on the Chr1. 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）

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Smyd2 SET and MYND domain containing 2 [Mus musculus (house mouse)]

Gene ID: 226830, updated on 31-Jan-2019

Summary

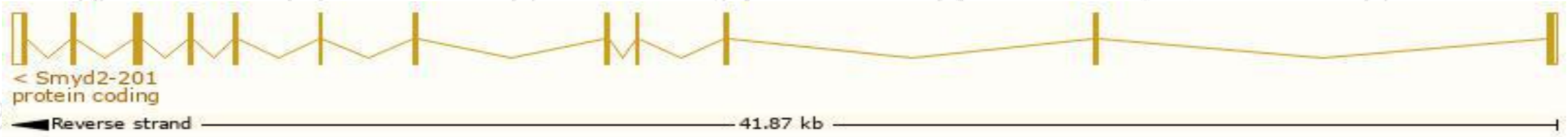
Official Symbol	Smyd2 provided by MGI
Official Full Name	SET and MYND domain containing 2 provided by MGI
Primary source	MGI:MGI:1915889
See related	Ensembl:ENSMUSG00000026603
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1110020E07Rik, 4930402C15, KMT3C, Zmynd14
Expression	Ubiquitous expression in ovary adult (RPKM 31.3), heart adult (RPKM 30.2) and 28 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

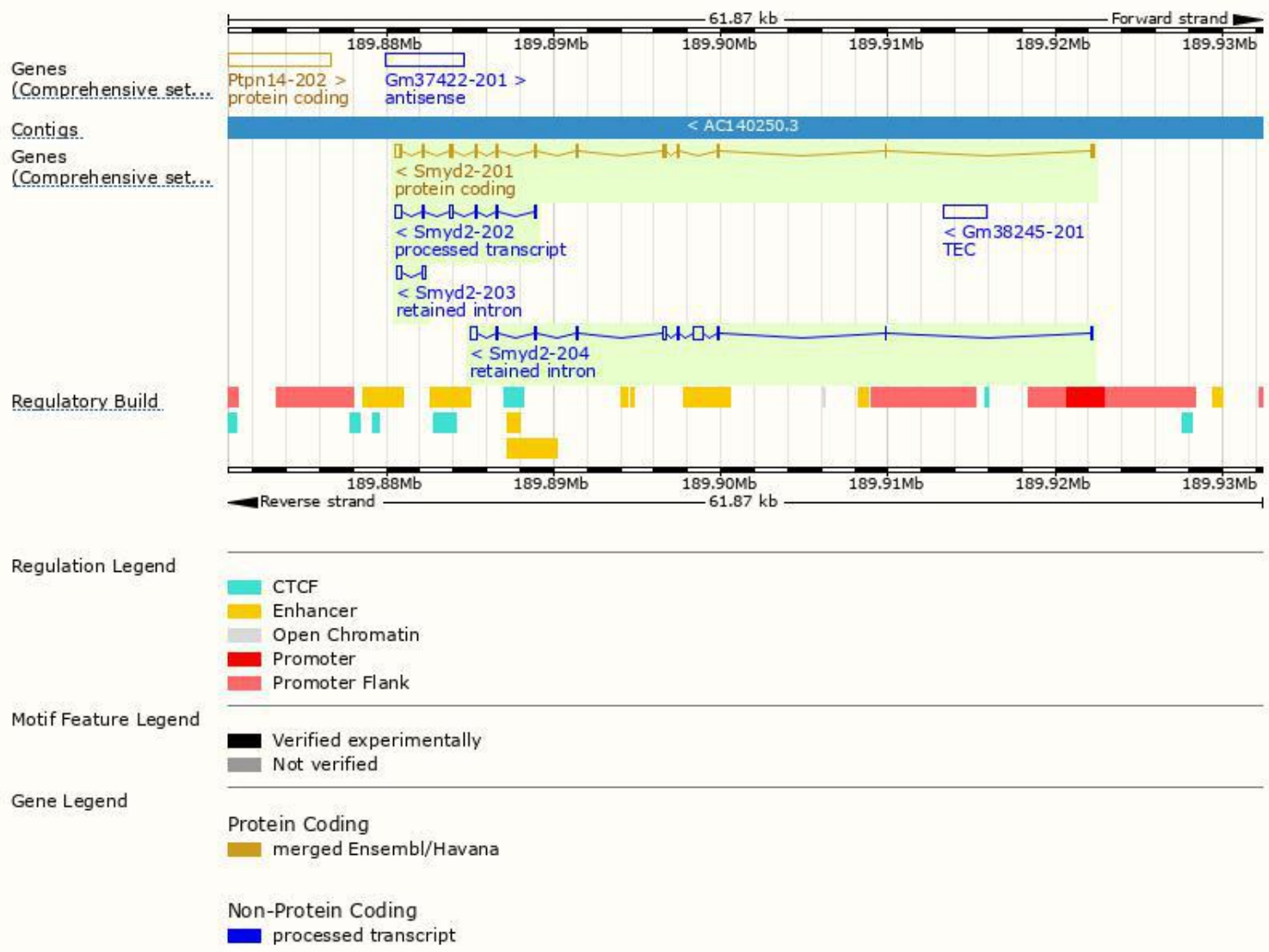
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Smyd2-201	ENSMUST00000027897.7	1680	433aa	Protein coding	CCDS35821	Q8R5A0	TSL:1 GENCODE basic APPRIS P1
Smyd2-202	ENSMUST00000130804.7	1077	No protein	Processed transcript	-	-	TSL:1
Smyd2-204	ENSMUST00000144452.1	1708	No protein	Retained intron	-	-	TSL:1
Smyd2-203	ENSMUST00000132289.1	489	No protein	Retained intron	-	-	TSL:1

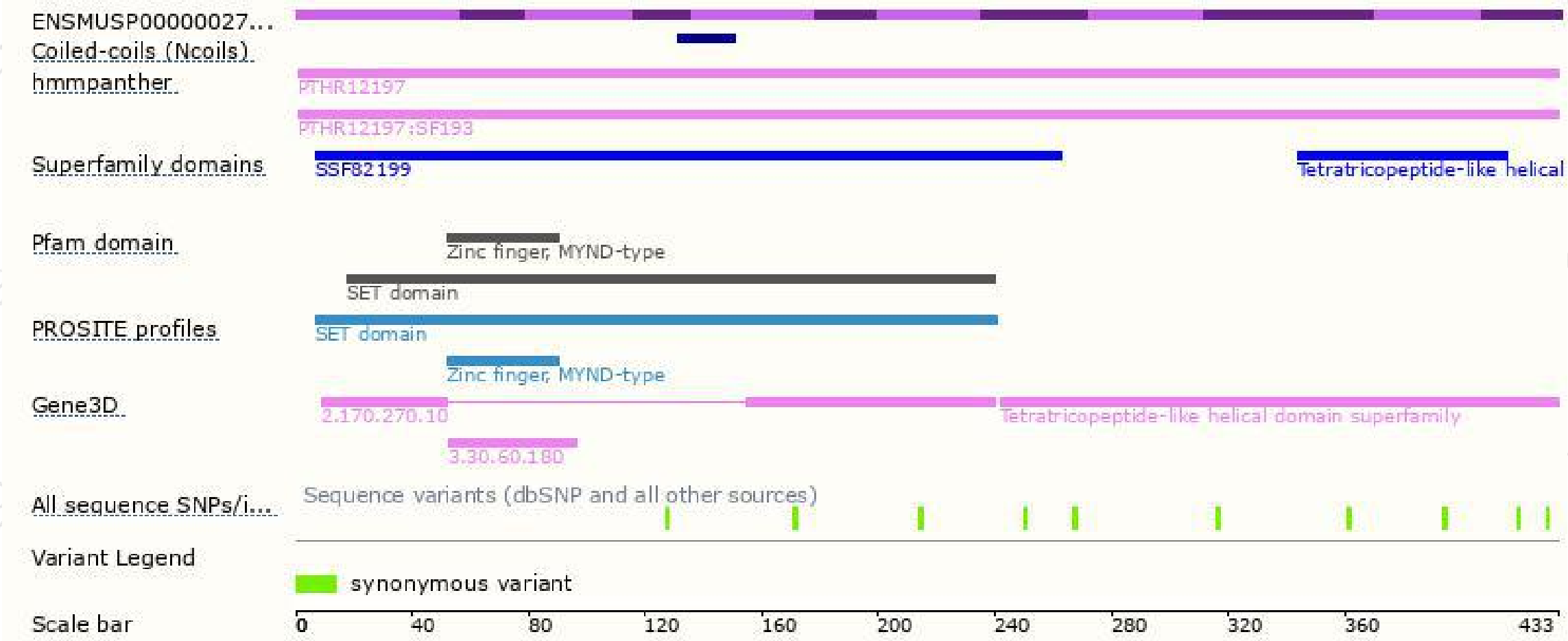
The strategy is based on the design of *Smyd2-201* transcript,The transcription is shown below



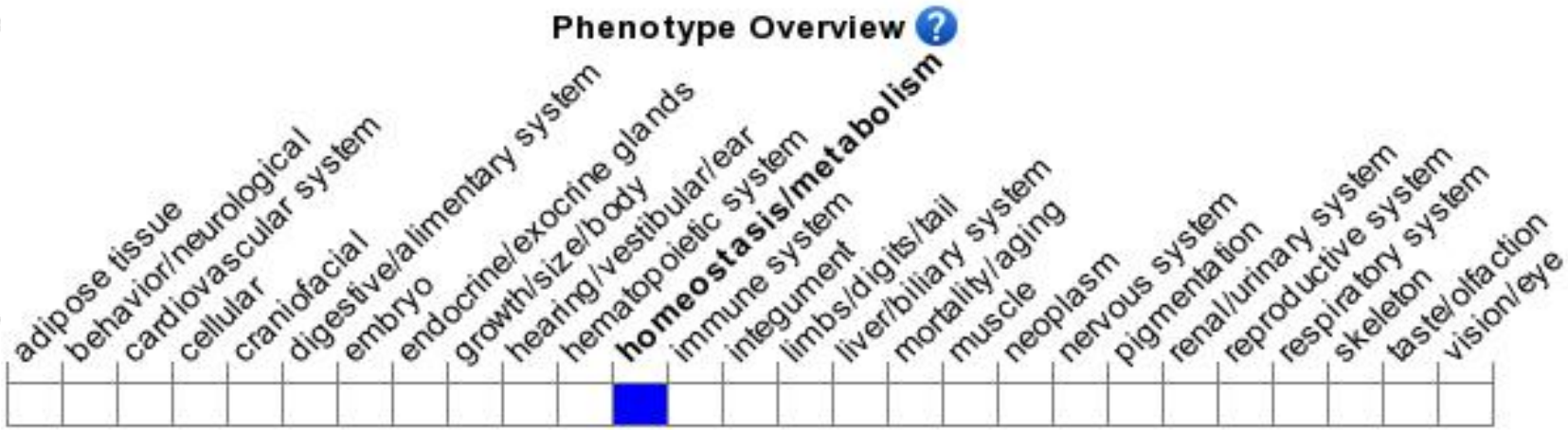
Genomic location distribution



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 targeted allele exhibit increased circulating total and LDL cholesterol levels and decreased circulating sodium and chloride levels.

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

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