Fabp1 Cas9-CKO *Strategy*

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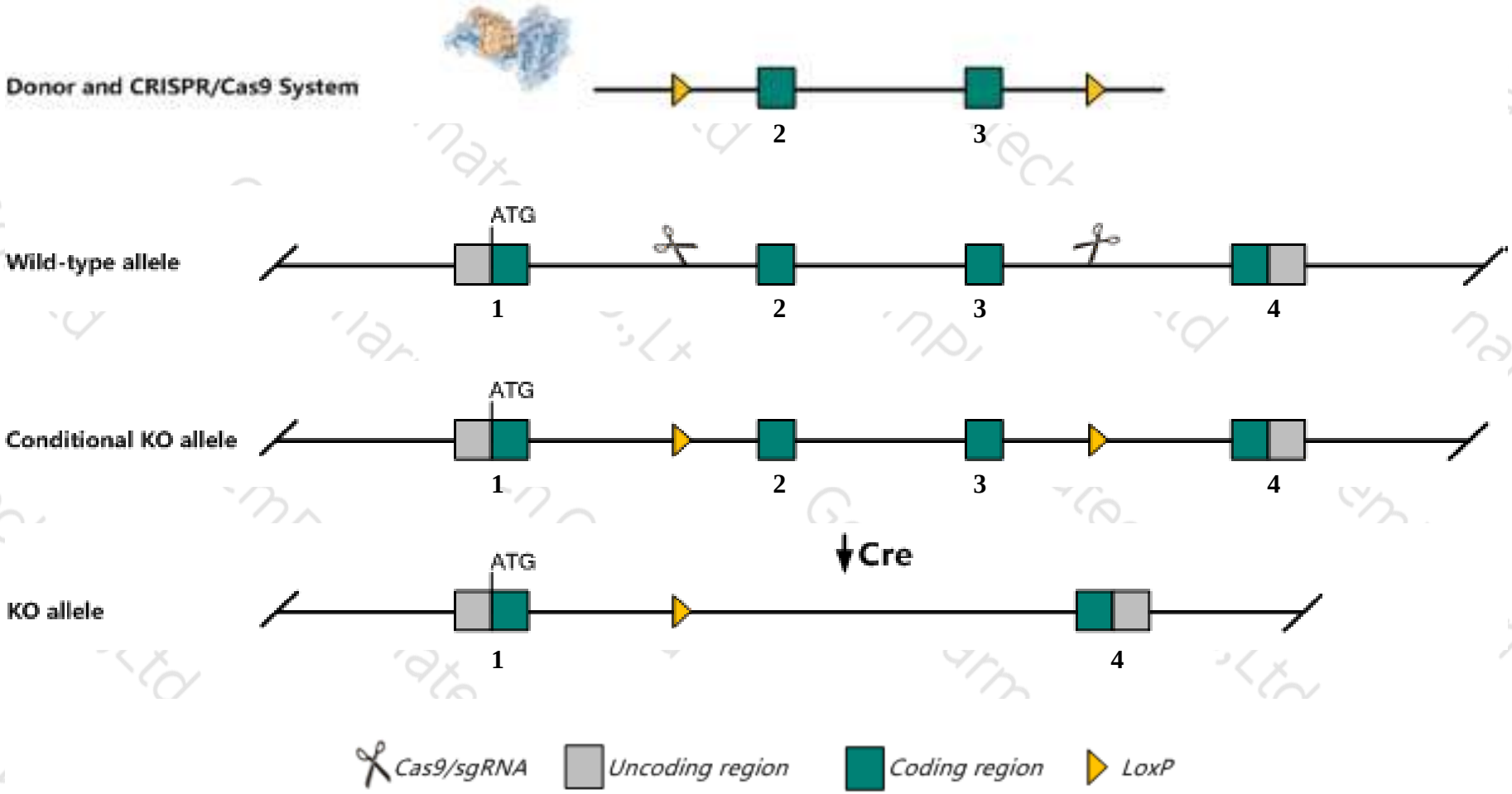
Project Overview

Project Name	***Fabp1***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fabp1* gene. The schematic diagram is as follows:



Technical routes

- The *Fabp1* gene has 1 transcript. According to the structure of *Fabp1* gene, exon2-exon3 of *Fabp1-201* (ENSMUST00000067492.7) transcript is recommended as the knockout region.The region contains 266bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Fabp1* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was 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 disruption in this gene do not show any obvious morphological, behavioral, or reproductive abnormalities. Changes may be observed in lipid composition, lipid binding, or levels of lipid binding proteins.

➢ The *Fabp1* gene is located on the Chr6. 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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Fabp1 fatty acid binding protein 1, liver [*Mus musculus* (house mouse)]

Gene ID: 14080, updated on 12-Aug-2019

Summary

Official Symbol	Fabp1 provided by MGI
Official Full Name	fatty acid binding protein 1, liver provided by MGI
Primary source	MGI:MGI:95479
See related	Ensembl:ENSMUSG00000054422
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	Fabpl; L-FABP
Expression	Biased expression in liver E18 (RPKM 1800.2), liver adult (RPKM 642.4) and 1 other tissue See more
Orthologs	human all

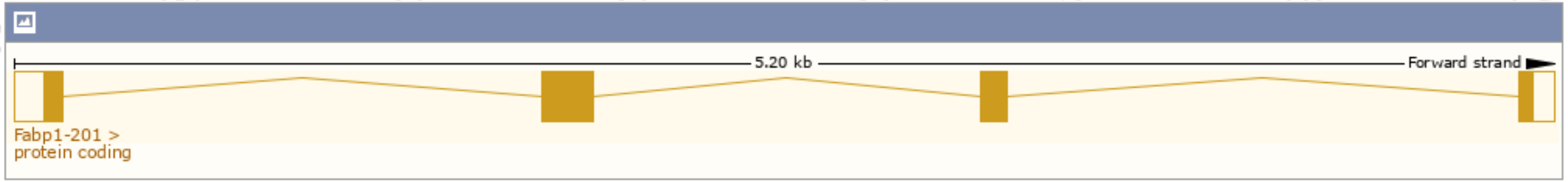
Transcript information（Ensembl）

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

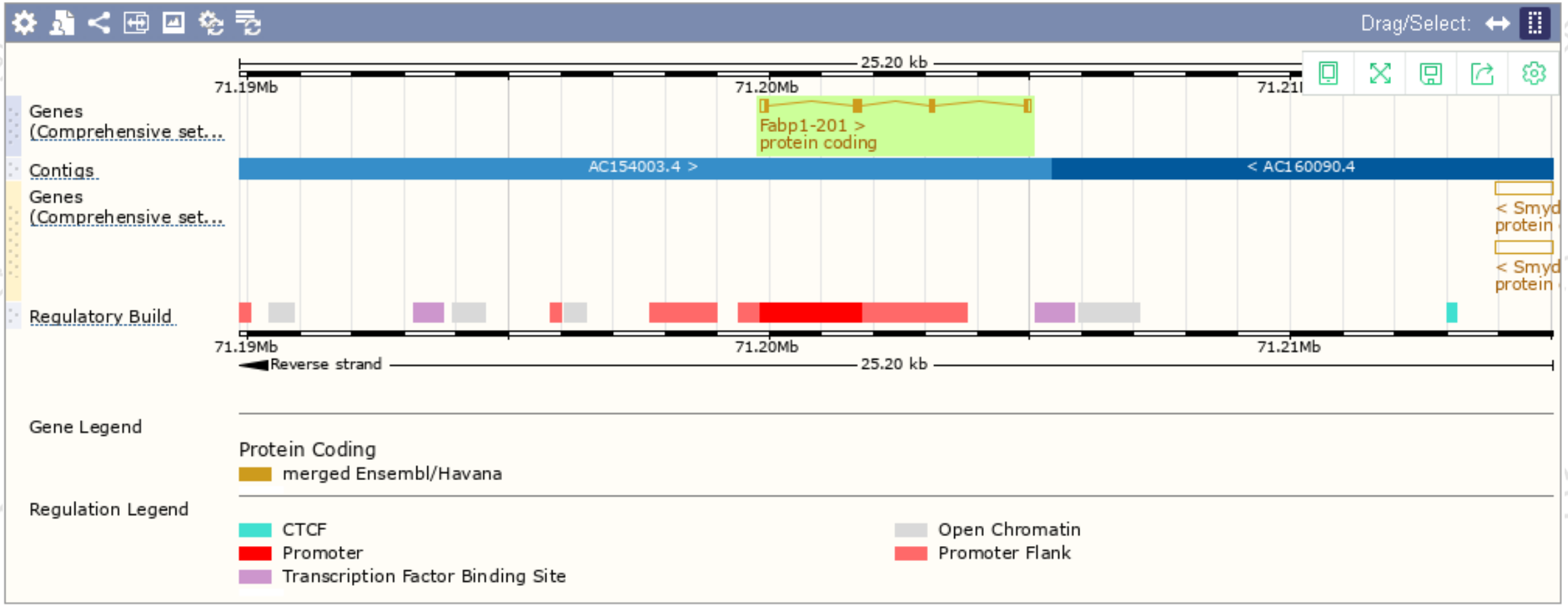
Show/hide columns (1 hidden) Filter

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fabp1-201	ENSMUST00000067492.7	553	127aa	Protein coding	CCDS20226	P12710 Q3V2F7	TSL:1 GENCODE basic APPRIS P1

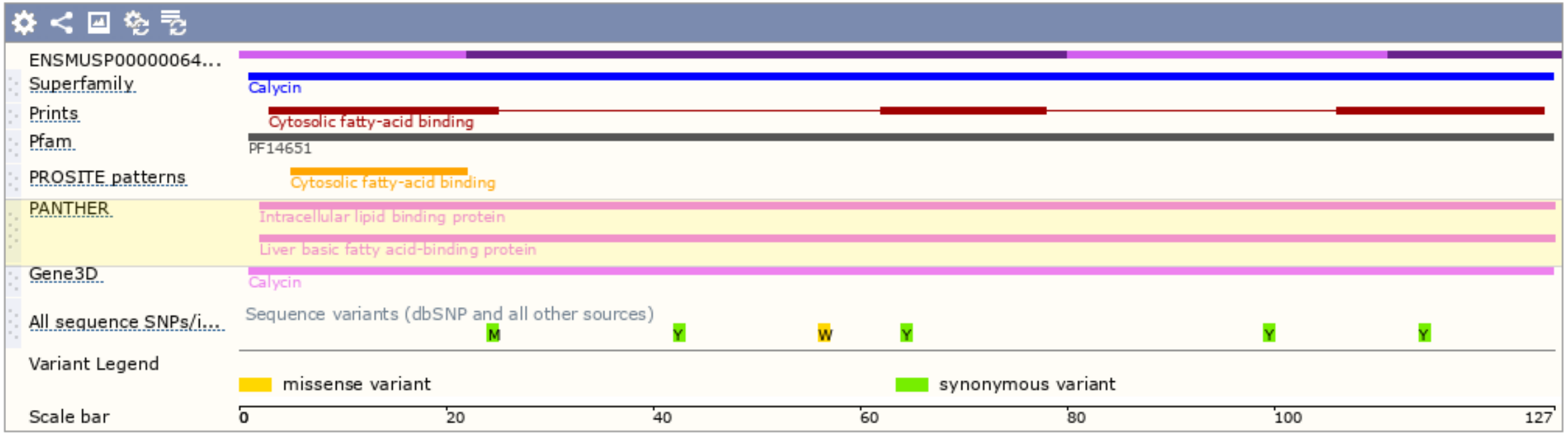
The strategy is based on the design of *Fabp1-201* transcript,The transcription is shown below:



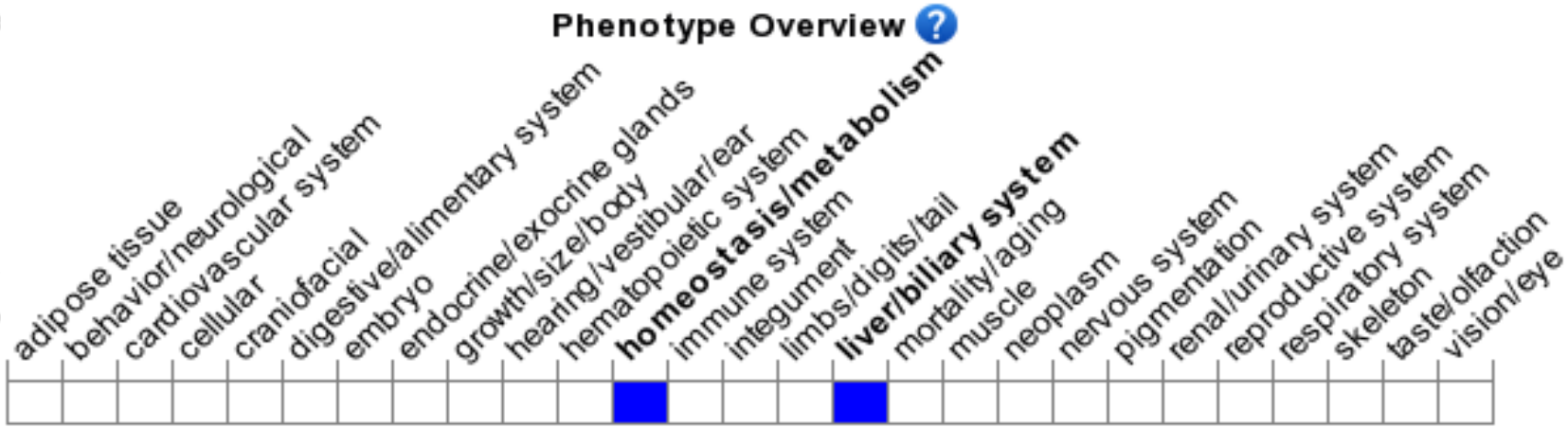
Genomic location distribution



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 disruption in this gene do not show any obvious morphological, behavioral, or reproductive abnormalities. Changes may be observed in lipid composition, lipid binding, or levels of lipid binding proteins.

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

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