Pla2g12b Cas9-CKO Strategy

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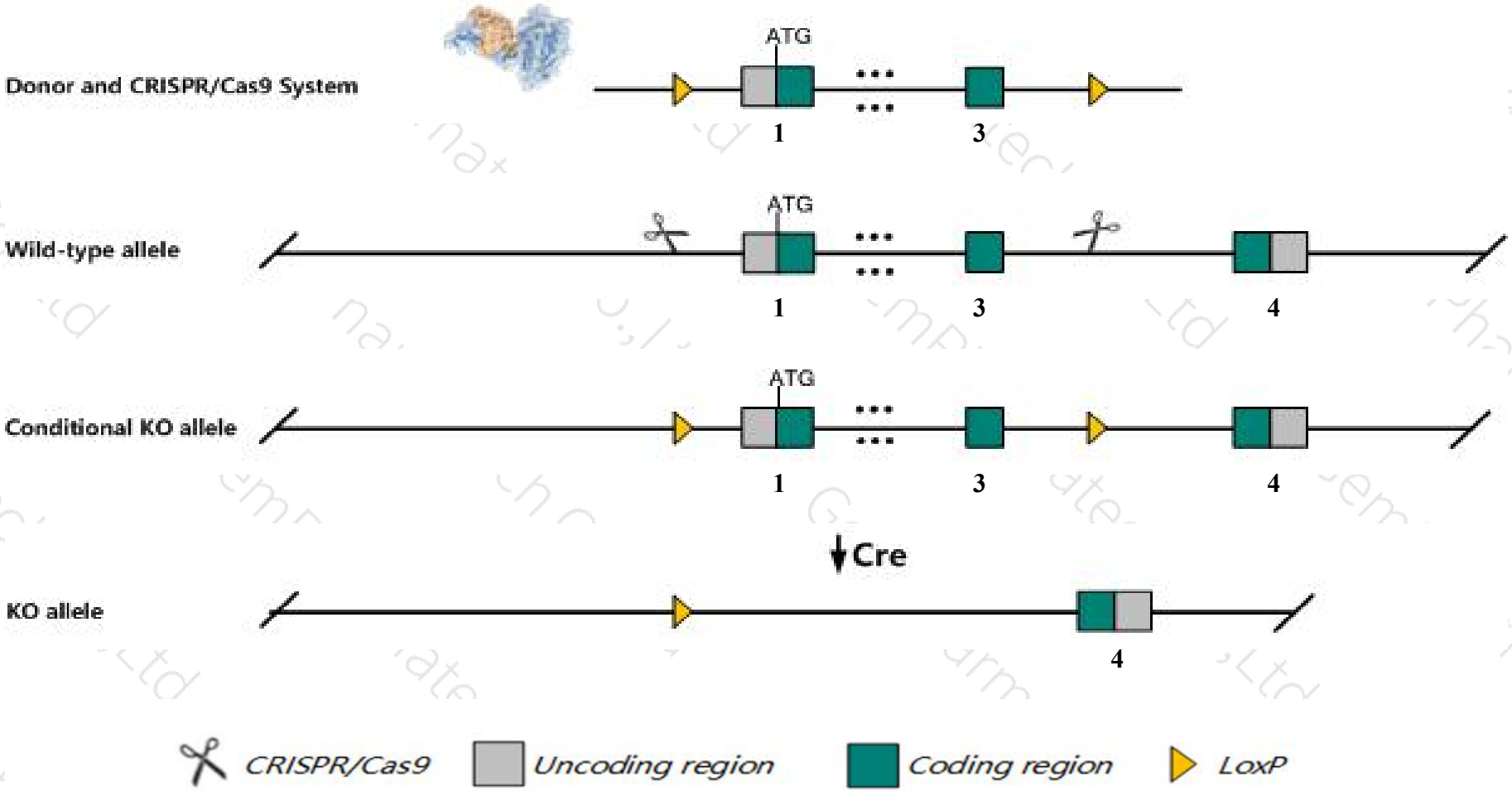
Project Overview

Project Name	***Pla2g12b***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Pla2g12b* gene. The schematic diagram is as follows:



Technical routes

- The *Pla2g12b* gene has 2 transcripts. According to the structure of *Pla2g12b* gene, exon1-exon3 of *Pla2g12b-201* (ENSMUST00000009790.13) transcript is recommended as the knockout region.The region contains start codon ATG. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Pla2g12b* 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 an enu-induced mutation exhibit a reduction in serum total and hdl cholesterol levels, decreased serum triglyceride levels, and hepatic steatosis.
- The *Pla2g12b* gene is located on the Chr10. 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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Pla2g12b phospholipase A2, group XIIB [Mus musculus (house mouse)]

Gene ID: 69836, updated on 13-Mar-2020

Summary

Official Symbol	Pla2g12b provided by MGI
Official Full Name	phospholipase A2, group XIIB provided by MGI
Primary source	MGI:MGI:1917086
See related	Ensembl:ENSMUSG00000009646
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	2010002E04Rik, Fksg71, Pla2g13, hlb218
Expression	Biased expression in placenta adult (RPKM 23.7), large intestine adult (RPKM 22.3) and 7 other tissues See more
Orthologs	human all

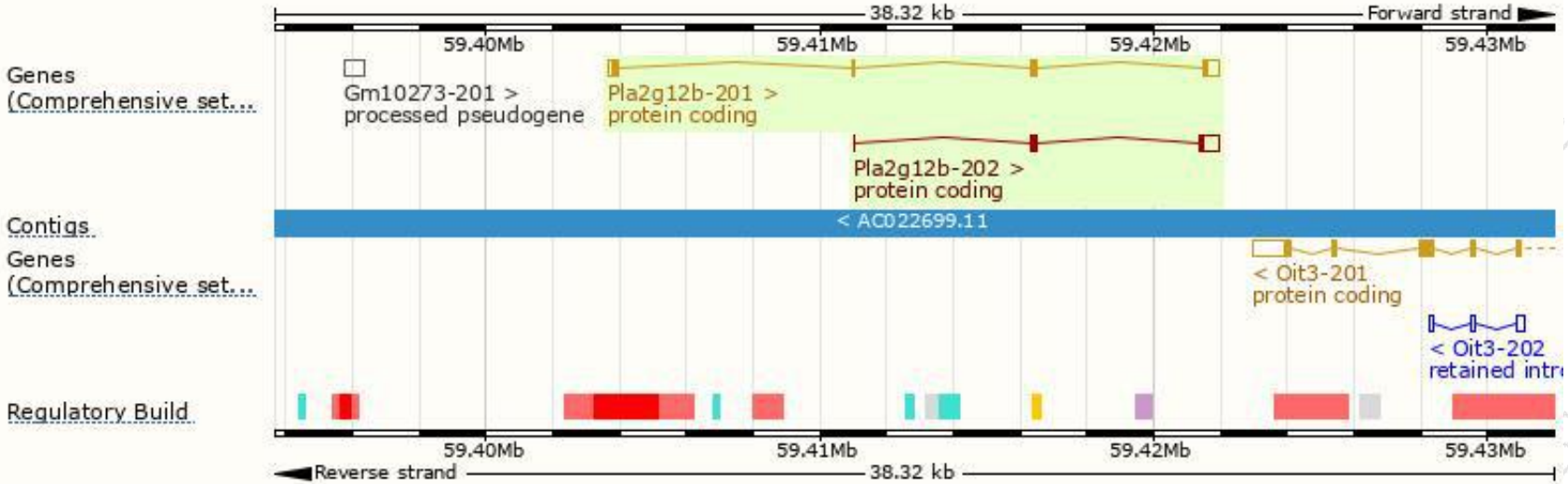
Transcript information（Ensembl）

The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pla2g12b-201	ENSMUST00000009790.13	1091	195aa	Protein coding	CCDS23864	Q8VC81	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1
Pla2g12b-202	ENSMUST00000162643.1	804	101aa	Protein coding	-	F6UTM6	CDS 5' incomplete TSL:2

The strategy is based on the design of *Pla2g12b-201* transcript,the transcription is shown below:

Genomic location distribution



Regulation Legend

- CTCF
- Enhancer
- Open Chromatin
- Promoter
- Promoter Flank
- Transcription Factor Binding Site

Gene Legend

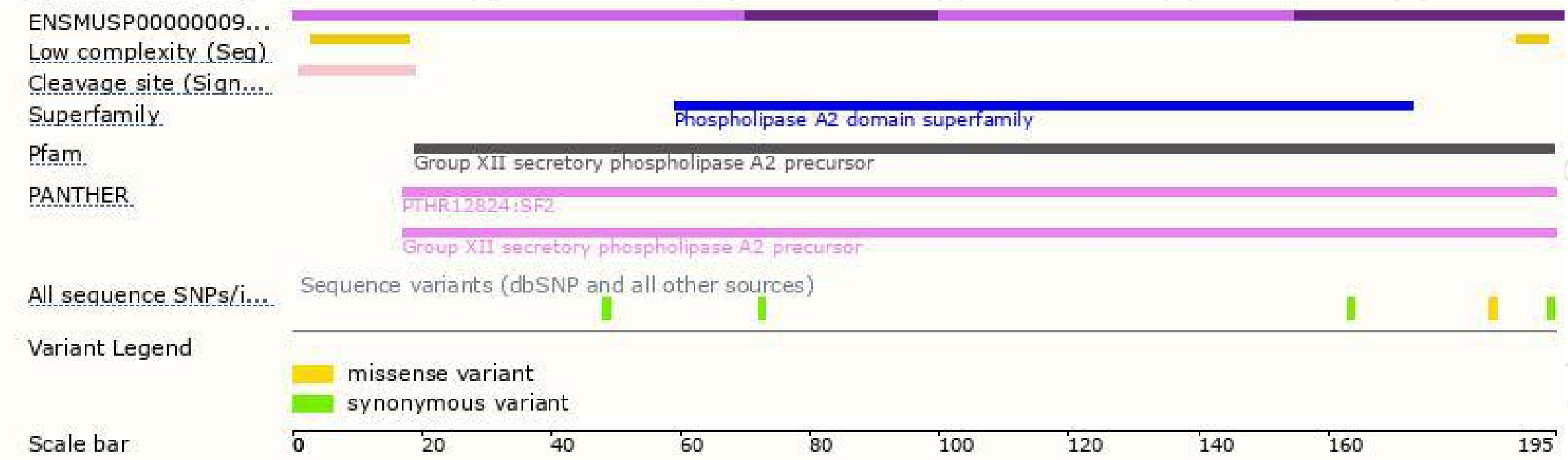
Protein Coding

- merged Ensembl/Havana
- Ensembl protein coding

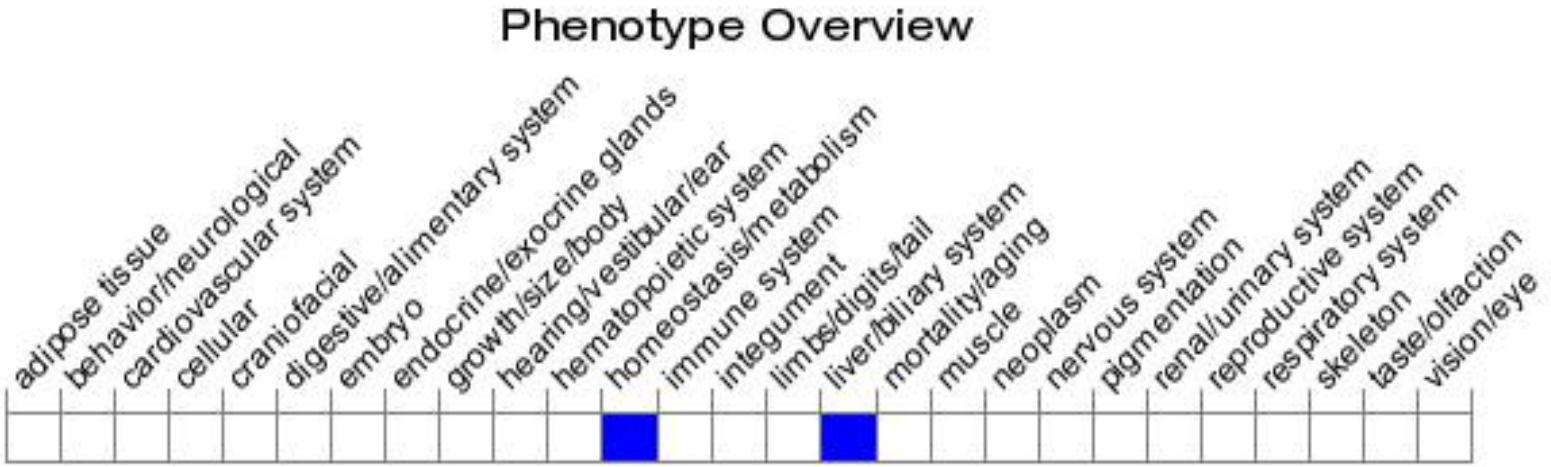
Non-Protein Coding

- pseudogene
- processed transcript

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 an ENU-induced mutation exhibit a reduction in serum total and HDL cholesterol levels, decreased serum triglyceride levels, and hepatic steatosis.

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

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