

Plcg2 Cas9-KO Strategy

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Project Overview

Project Name

Plcg2

Project type

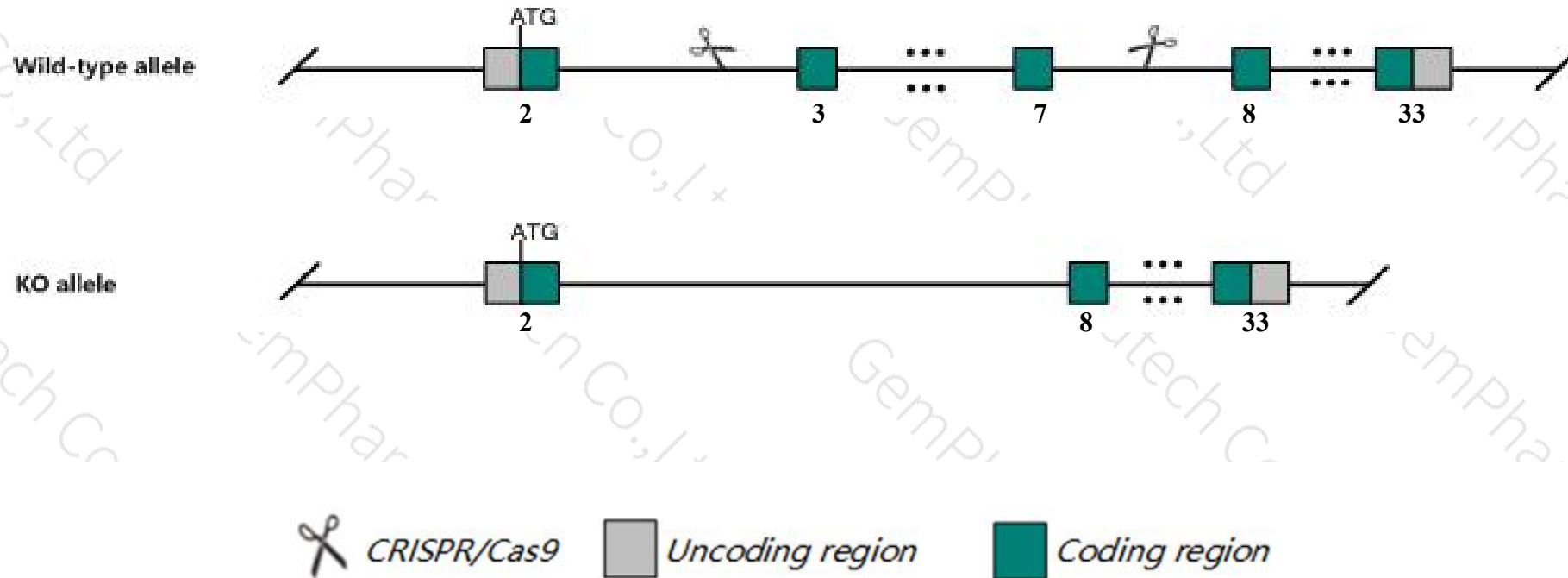
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Plcg2* gene. The schematic diagram is as follows:



- The *Plcg2* gene has 1 transcript. According to the structure of *Plcg2* gene, exon3-exon7 of *Plcg2*-201 (ENSMUST00000081232.8) transcript is recommended as the knockout region. The region contains 455bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Plcg2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygotes for some null alleles show decreased B cell and impaired NK cell function. Other homozygous null alleles show aberrant separation of blood and lymphatic vessels.
- The *Plcg2* gene is located on the Chr8. 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)

Plcg2 phospholipase C, gamma 2 [*Mus musculus* (house mouse)]

Gene ID: 234779, updated on 10-Oct-2019

Summary

Official Symbol	Plcg2 provided by MGI
Official Full Name	phospholipase C, gamma 2 provided by MGI
Primary source	MGI:MGI:97616
See related	Ensembl:ENSMUSG00000034330
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<i>Mus musculus</i>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Plcg-2; PLCgamma2; PLC-gamma-2
Expression	Broad expression in spleen adult (RPKM 27.6), mammary gland adult (RPKM 7.2) and 19 other tissues See more
Orthologs	human all

Genomic context

Location: 8 E1; 8 64.26 cM

See Plcg2 in [Genome Data Viewer](#)

Exon count: 33

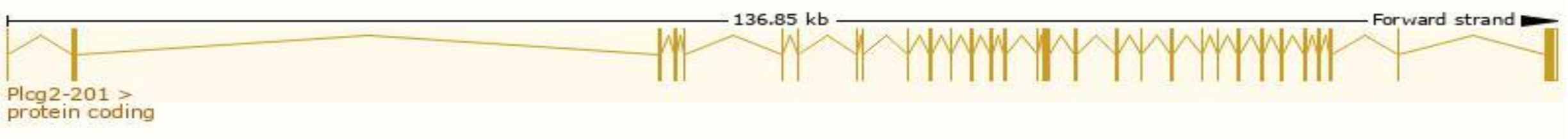
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	8	NC_000074.6 (117498275..117635142)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	8	NC_000074.5 (120022191..120159042)

Transcript information (Ensembl)

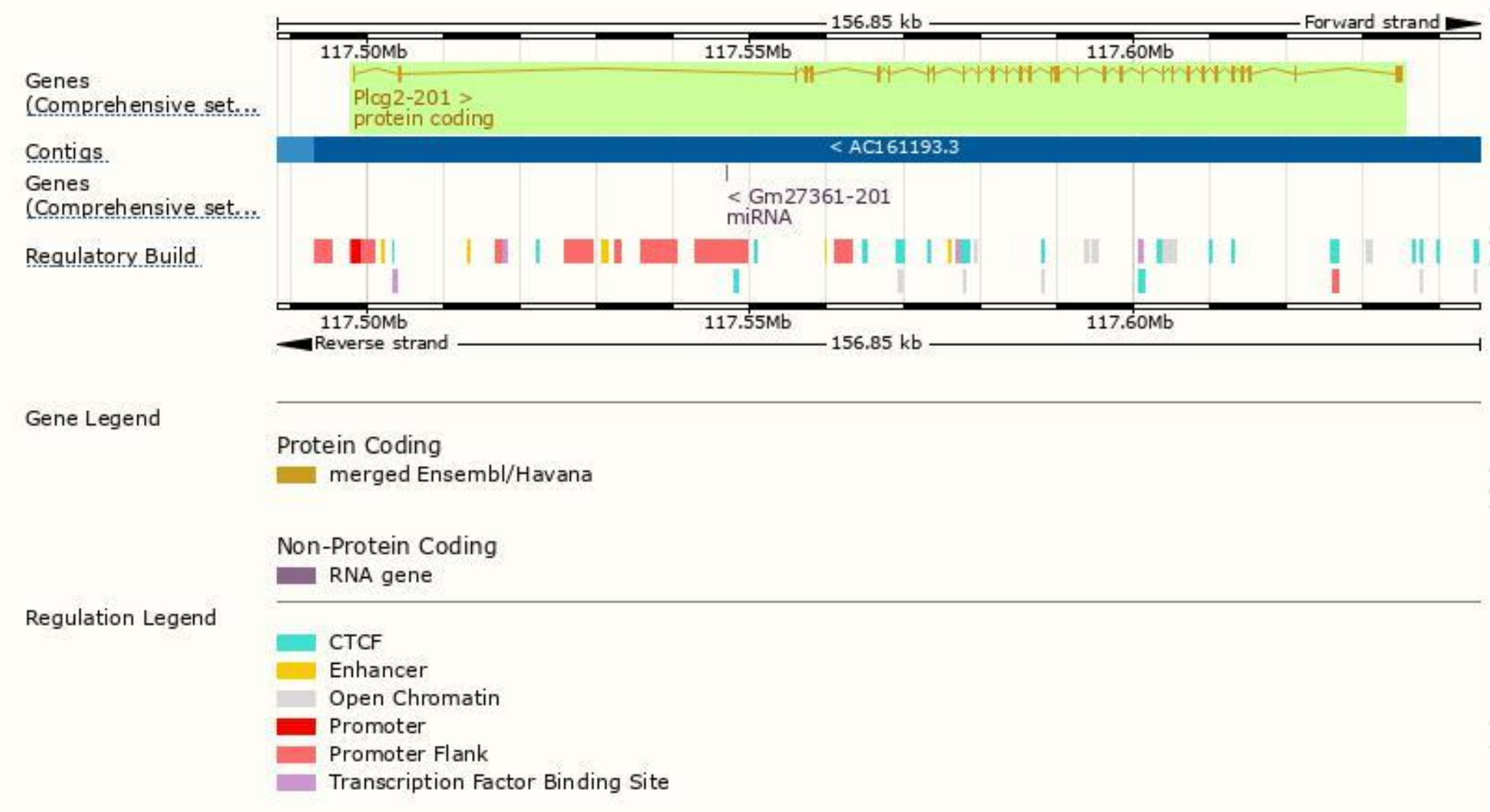
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Plcg2-201	ENSMUST00000081232.8	4298	1265aa	Protein coding	CCDS22700	Q8CIH5	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Plcg2-201* transcript, The transcription is shown below



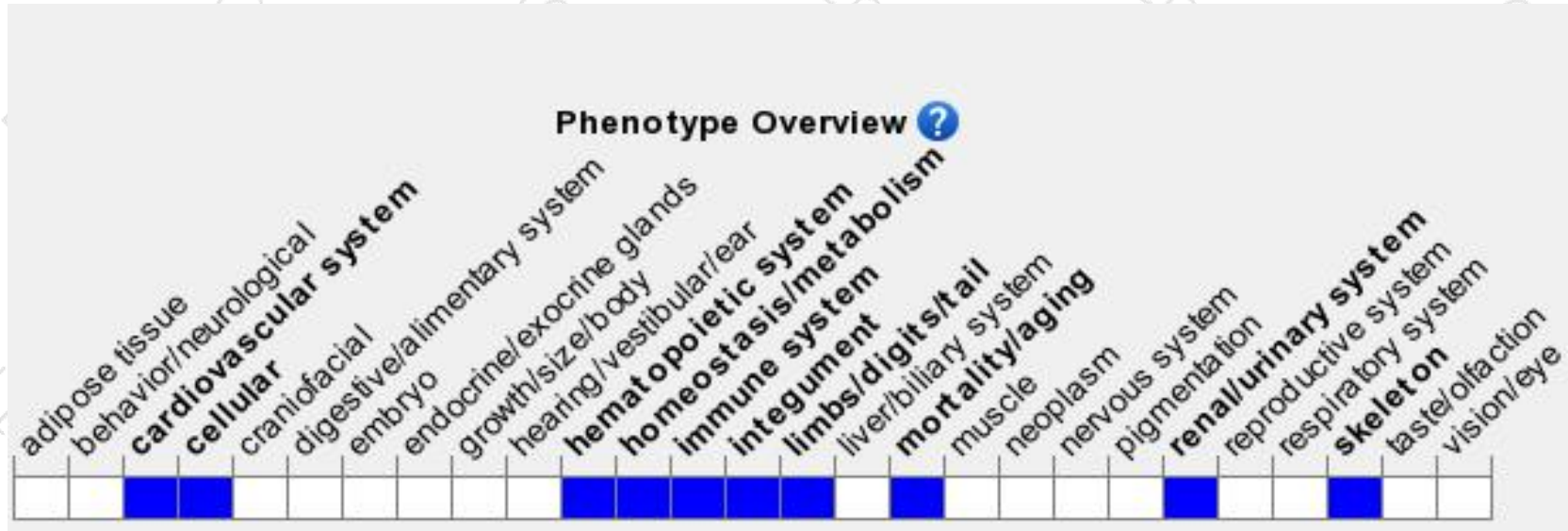
Genomic location distribution



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, Homozygotes for some null alleles show decreased B cell and impaired NK cell function. Other homozygous null alleles show aberrant separation of blood and lymphatic vessels.

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

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