

Plcl1 Cas9-KO Strategy

Designer: Xueting Zhang

Reviewer: Yanhua Shen

Date: 2020-1-19

Project Overview

Project Name

Plcl1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Plcl1* gene has 2 transcripts. According to the structure of *Plcl1* gene, exon2-exon4 of *Plcl1-201* (ENSMUST00000042986.9) transcript is recommended as the knockout region. The region contains 2755bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Plcl1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Homozygous null mutants display impaired motor coordination and decreased sensitivity to the sedative diazepam.
- The *Plcl1* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Plcl1 phospholipase C-like 1 [*Mus musculus* (house mouse)]

Gene ID: 227120, updated on 10-Dec-2019

Summary

Official Symbol Plcl1 provided by [MGI](#)
Official Full Name phospholipase C-like 1 provided by [MGI](#)
Primary source [MGI:MGI:3036262](#)
See related [Ensembl:ENSMUSG00000038349](#)
Gene type protein coding
RefSeq status PROVISIONAL
Organism [Mus musculus](#)
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as PLCE; PLCL; PRIP; PLC-L; PLDL1; PLC-L1; PRIP-1; C230017K02Rik
Expression Broad expression in bladder adult (RPKM 3.1), cerebellum adult (RPKM 3.0) and 24 other tissues [See more](#)
Orthologs [human](#) [all](#)

Genomic context

Location: 1; 1 C1.2

See Plcl1 in [Genome Data Viewer](#)

Exon count: 6

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	1	NC_000067.6 (55405946..55754285)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	1	NC_000067.5 (55462790..55811129)

Transcript information (Ensembl)

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Plcl1-201	ENSMUST00000042986.9	6580	1096aa	Protein coding	CCDS48263	Q3USB7	TSL:1 GENCODE basic APPRIS P1
Plcl1-202	ENSMUST00000187059.1	402	No protein	lncRNA	-	-	TSL:2

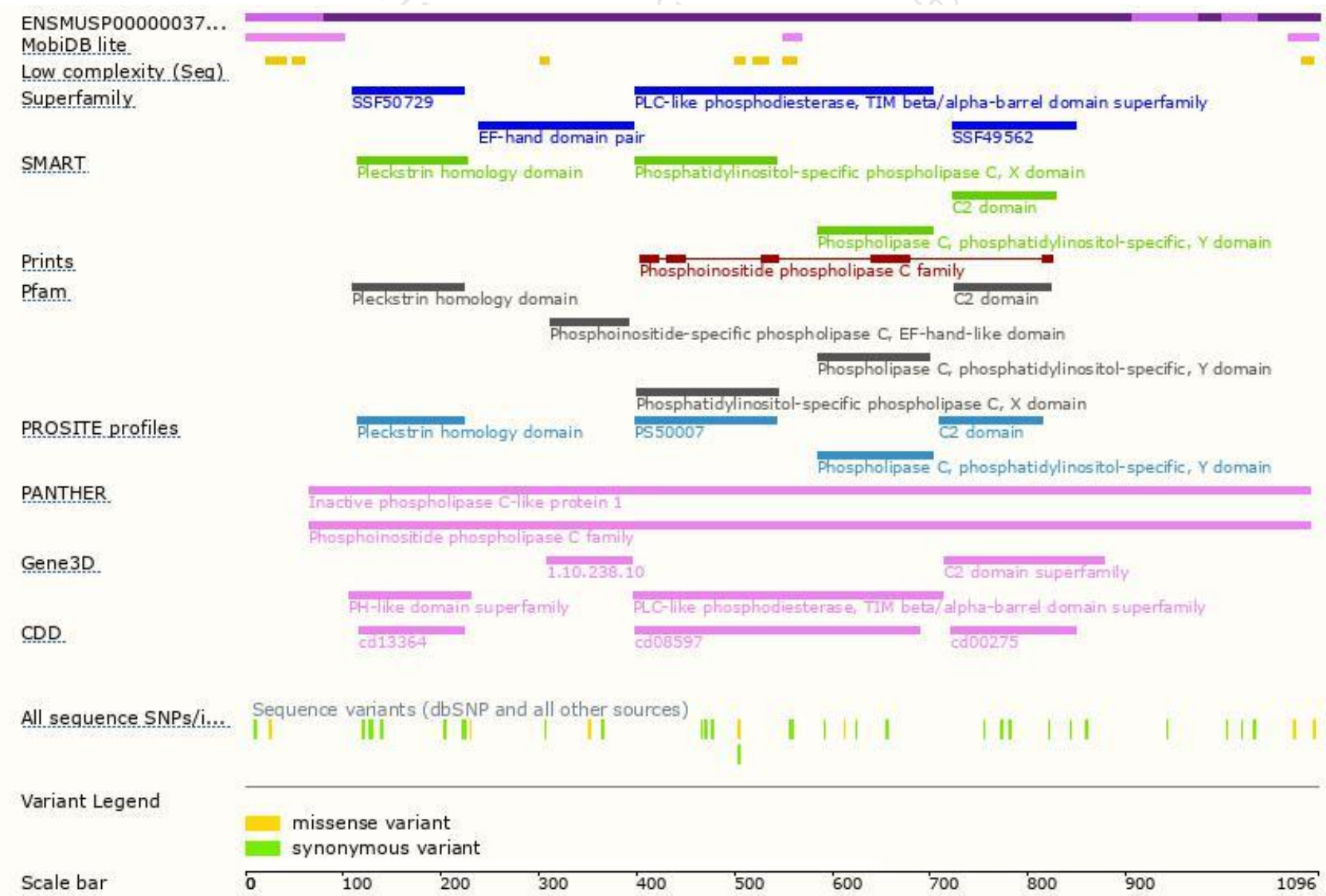
The strategy is based on the design of *Plcl1-201* transcript,The transcription is shown below



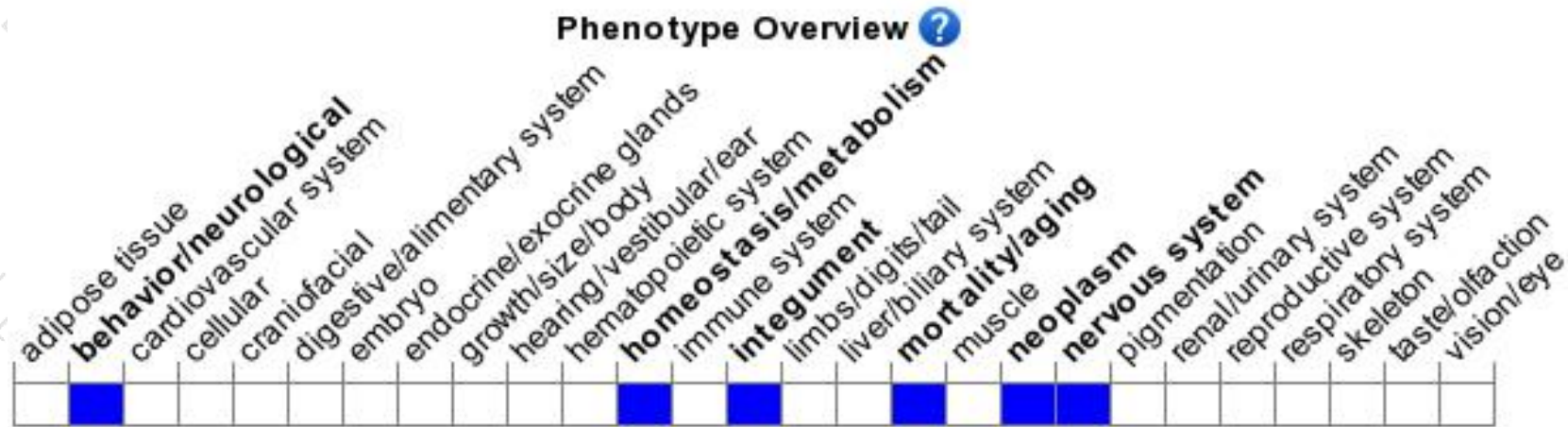
Genomic location distribution



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, Homozygous null mutants display impaired motor coordination and decreased sensitivity to the sedative diazepam.

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

Tel: 400-9660890

