

Pld6 Cas9-KO Strategy

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Project Overview



Project Name

Project type

Strain background

Cas9-KO

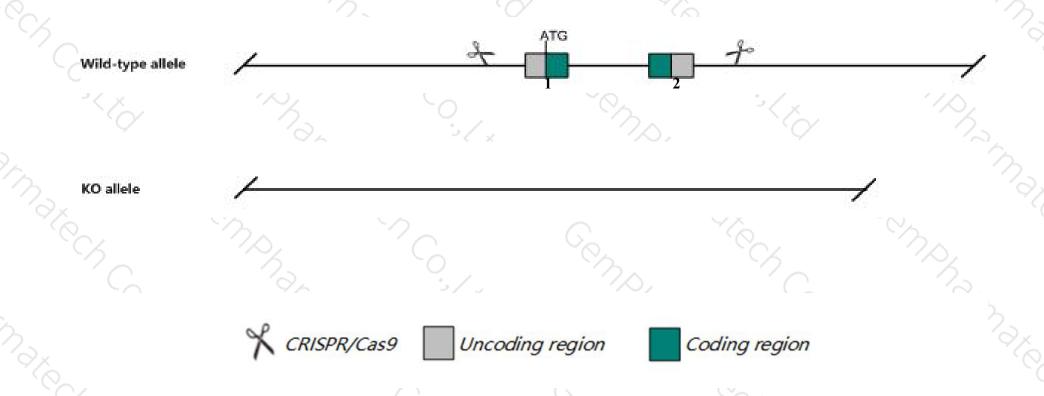
Pld6

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pld6* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Pld6* gene has 2 transcripts. According to the structure of *Pld6* gene, exon1-exon2 of *Pld6-202* (ENSMUST00000125307.1) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Pld6* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice



- > According to the existing MGI data, Mice homozygous for a null mutation display male infertility with arrest of male meiosis and mitochondrial abnormalities.
- The knockout region is near to the N-terminal of Gm12264 gene, this strategy may influence the regulatory function of the N-terminal of Gm12264 gene.
- > The *Pld6* gene is located on the Chr11. 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)



Pld6 phospholipase D family, member 6 [Mus musculus (house mouse)]

Gene ID: 194908, updated on 3-Feb-2019

Summary

☆ ?

Official Symbol Pld6 provided by MGI

Official Full Name phospholipase D family, member 6 provided by MGI

Primary source MGI:MGI:2687283

See related Ensembl: ENSMUSG00000043648

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 4933433K01Rik, Gm10, Zuc, mZuc, mitoPLD

Expression Biased expression in testis adult (RPKM 34.0), stomach adult (RPKM 4.9) and 3 other tissuesSee more

Orthologs <u>human</u> all

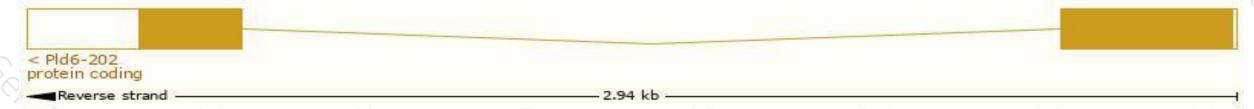
Transcript information (Ensembl)



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

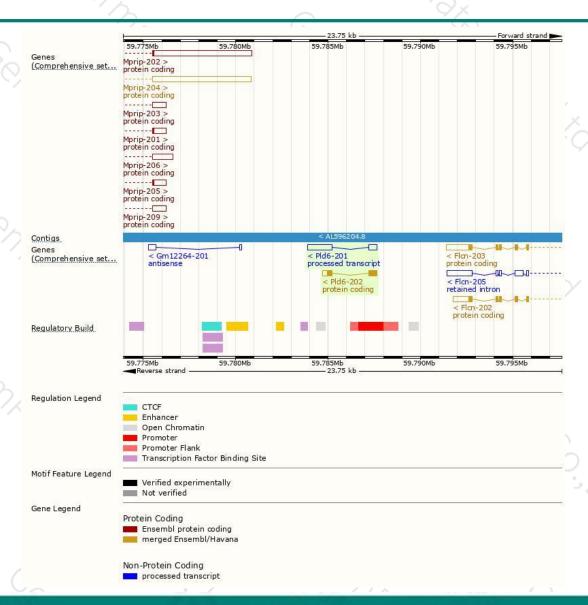
Name	Transcript ID	bp	Protein	Biotype	ccds	UniProt	Flags
PId6-202	ENSMUST00000125307.1	948	221aa	Protein coding	CCDS70199	Q5SWZ9	TSL:1 GENCODE basic APPRIS P1
PId6-201	ENSMUST00000051493.7	1753	No protein	Processed transcript	5		TSL:1

The strategy is based on the design of *Pld6-202* transcript, The transcription is shown below



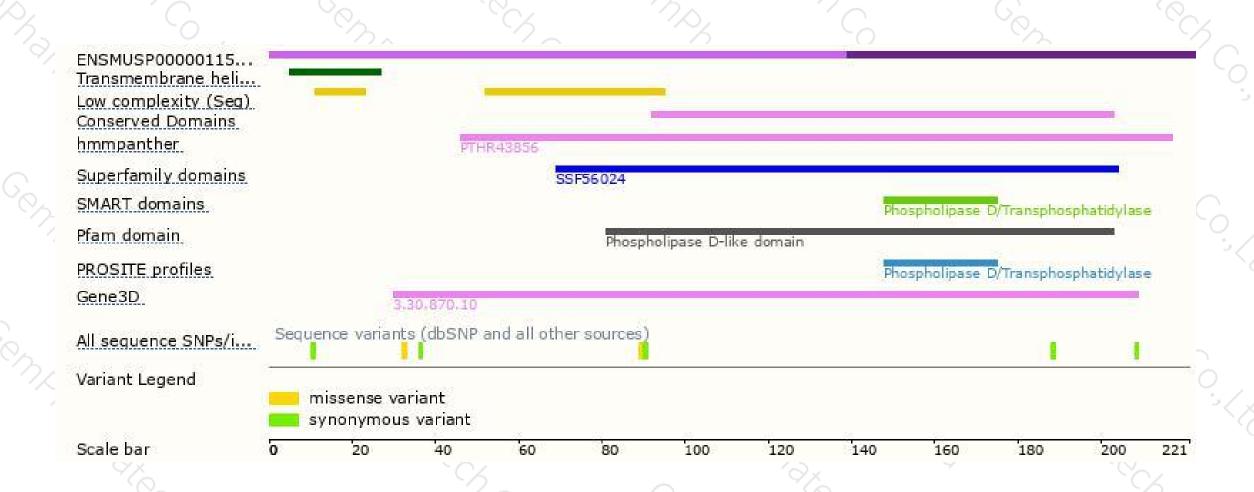
Genomic location distribution





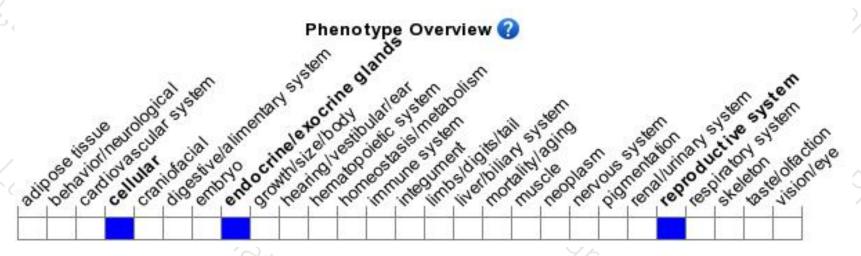
Protein domain





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 null mutation display male infertility with arrest of male meiosis and mitochondrial abnormalities.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





