

Pinlyp Cas9-KO Strategy

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Design Date:

Project Overview

Project Name

Pinlyp

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Pinlyp* gene has 1 transcript. According to the structure of *Pinlyp* gene, exon3-exon5 of *Pinlyp*-201 (ENSMUST00000011776.7) transcript is recommended as the knockout region. The region contains 452bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Pinlyp* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Male mice homozygous for a mutation are viable and show normal fertility.
- Pinlyp gene is less than 5kb away from Xrcc1, which may affect the activation of Pinlyp promoter.
- The *Pinlyp* gene is located on the Chr7. 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)

Pinlyp phospholipase A2 inhibitor and LY6/PLAUR domain containing [Mus musculus (house mouse)]

Gene ID: 641361, updated on 31-Jan-2019

Summary



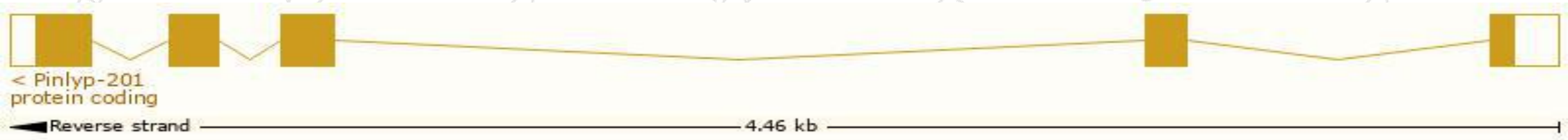
Official Symbol	Pinlyp provided by MGI
Official Full Name	phospholipase A2 inhibitor and LY6/PLAUR domain containing provided by MGI
Primary source	MGI:MGI:3615324
See related	Ensembl:ENSMUSG00000011632
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2310033E01Rik
Expression	Biased expression in stomach adult (RPKM 15.5), testis adult (RPKM 6.0) and 3 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

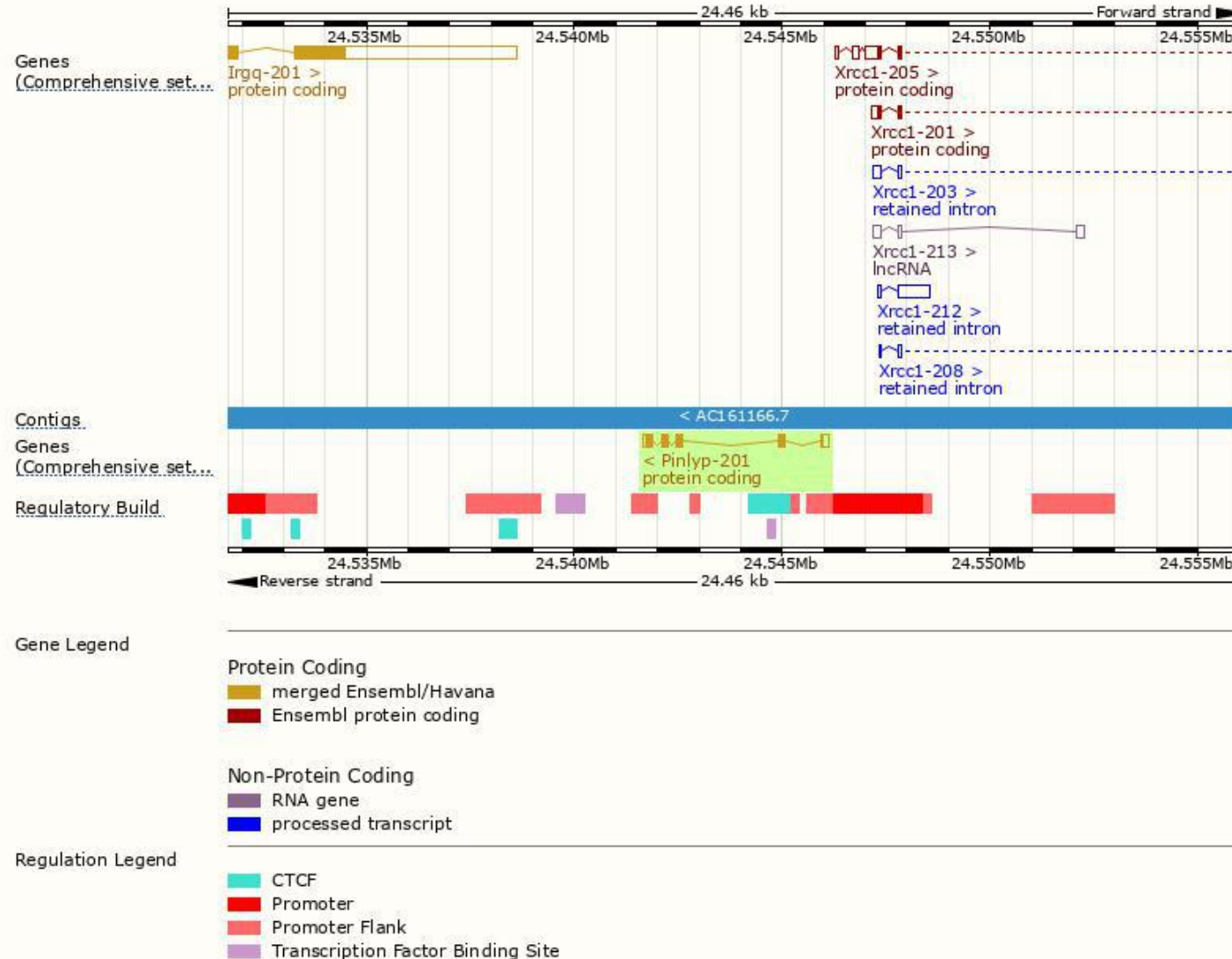
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pinlyp-201	ENSMUST00000011776.7	843	212aa	Protein coding	CCDS20954	B2RSX2 Q9CQD7	TSL:1 GENCODE basic APPRIS P1

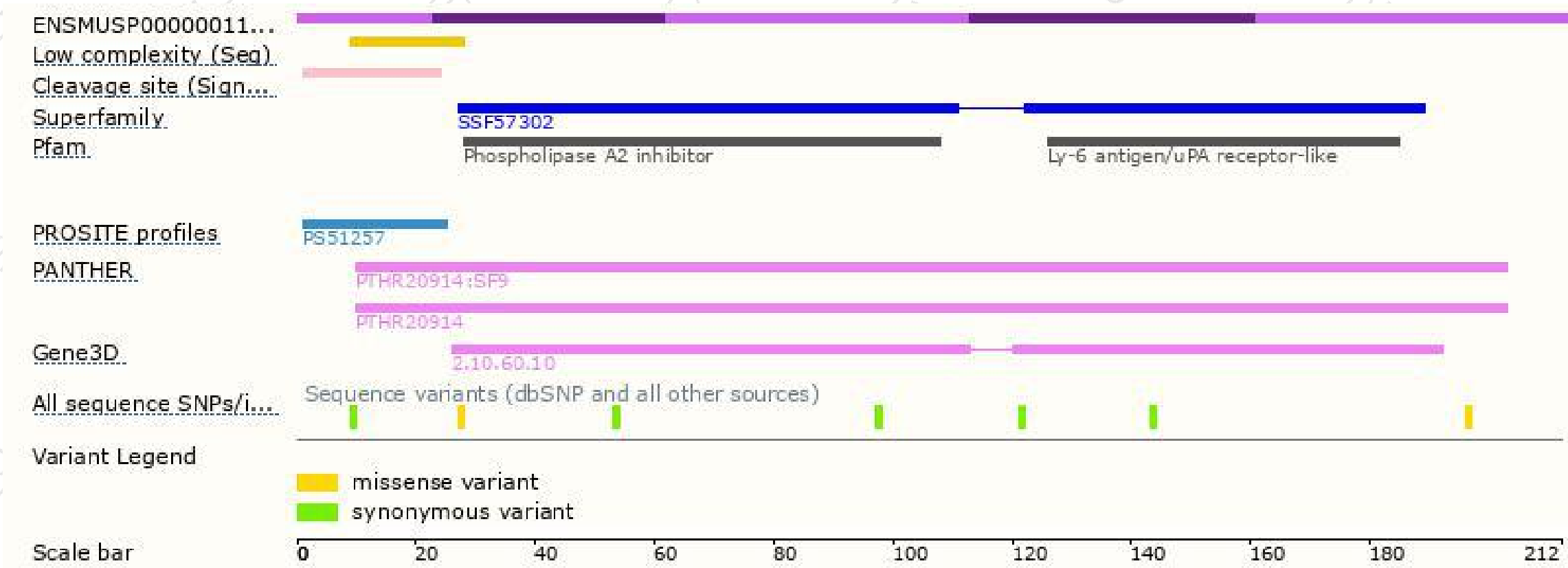
The strategy is based on the design of *Pinlyp-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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