

***Pard6g* Cas9-KO Strategy**

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Design Date: 2020-4-16

Project Overview

Project Name

Pard6g

Project type

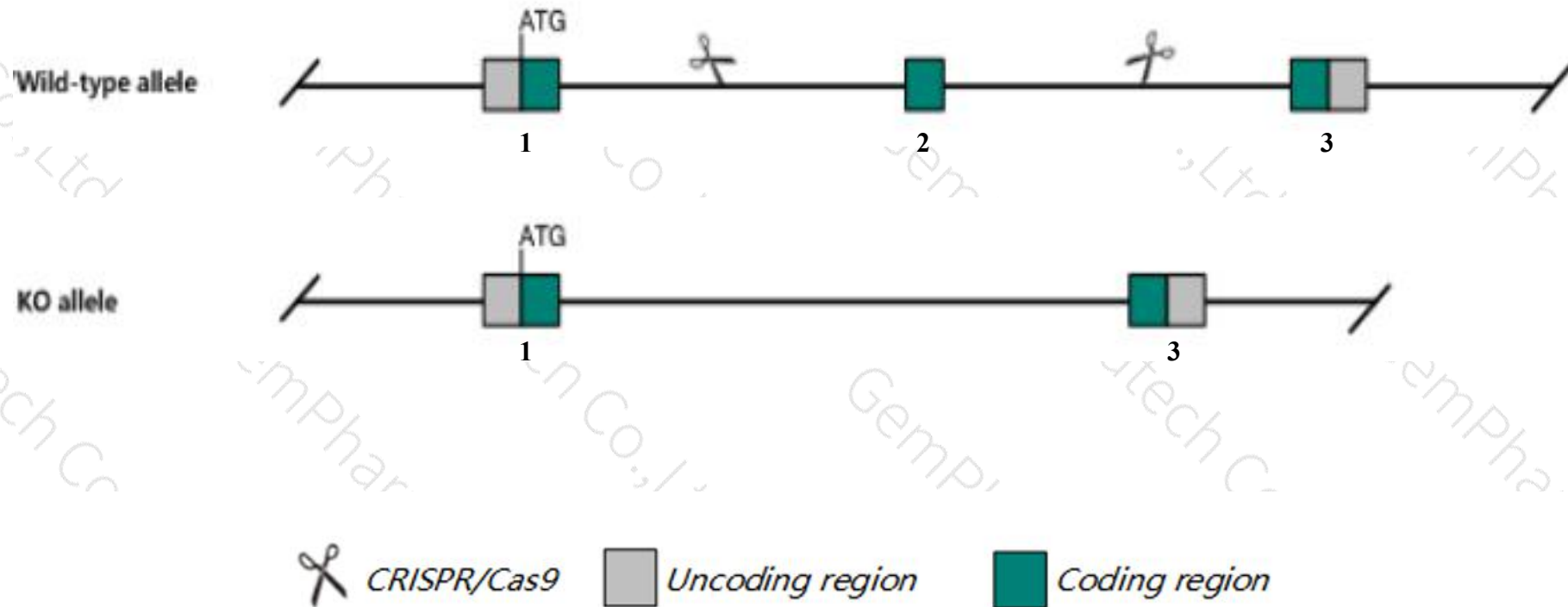
Cas9-KO

Strain background

C57BL/6J

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Pard6g* gene. The schematic diagram is as follows:



- The *Pard6g* gene has 1 transcript. According to the structure of *Pard6g* gene, exon2 of *Pard6g-201* (ENSMUST00000070219.8) transcript is recommended as the knockout region. The region contains 223bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Pard6g* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Pard6g* gene is located on the Chr18. 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)

Pard6g par-6 family cell polarity regulator gamma [Mus musculus (house mouse)]

Gene ID: 93737, updated on 13-Mar-2020

Summary



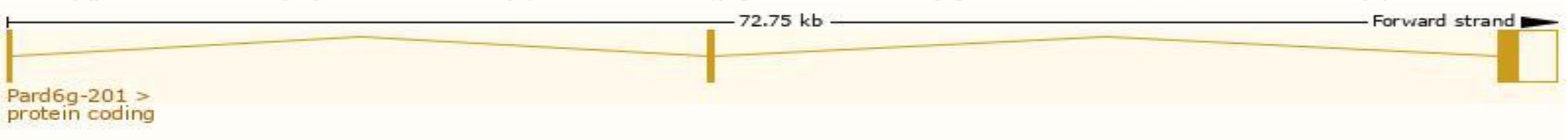
Official Symbol	Pard6g provided by MGI
Official Full Name	par-6 family cell polarity regulator gamma provided by MGI
Primary source	MGI:MGI:2135606
See related	Ensembl:ENSMUSG00000056214
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2410049N21Rik, Par6a
Expression	Broad expression in limb E14.5 (RPKM 8.4), CNS E11.5 (RPKM 8.1) and 23 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

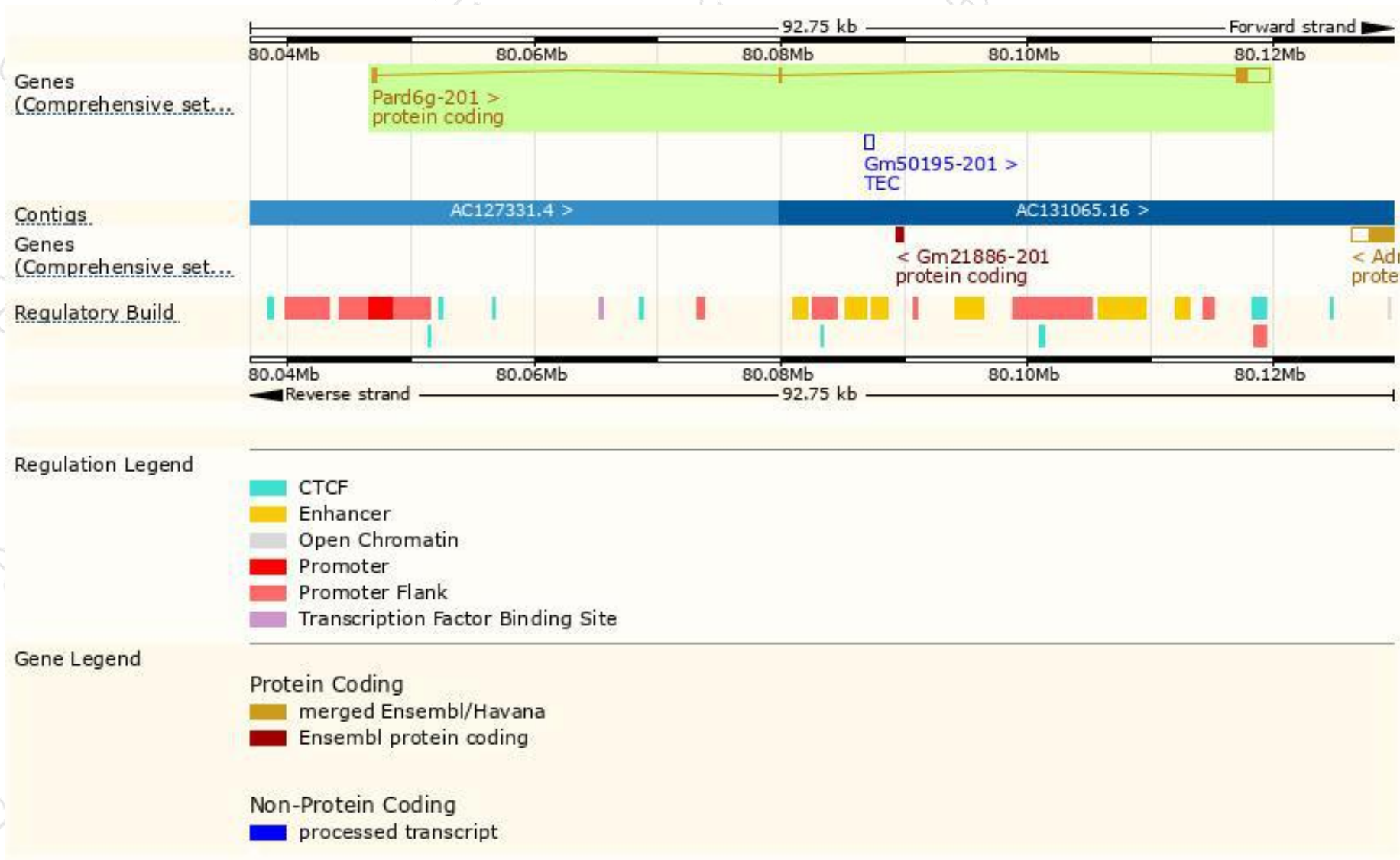
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pard6g-201	ENSMUST00000070219.8	3147	382aa	Protein coding	CCDS29363	Q9JK84	TSL:1 GENCODE basic APPRIS P1

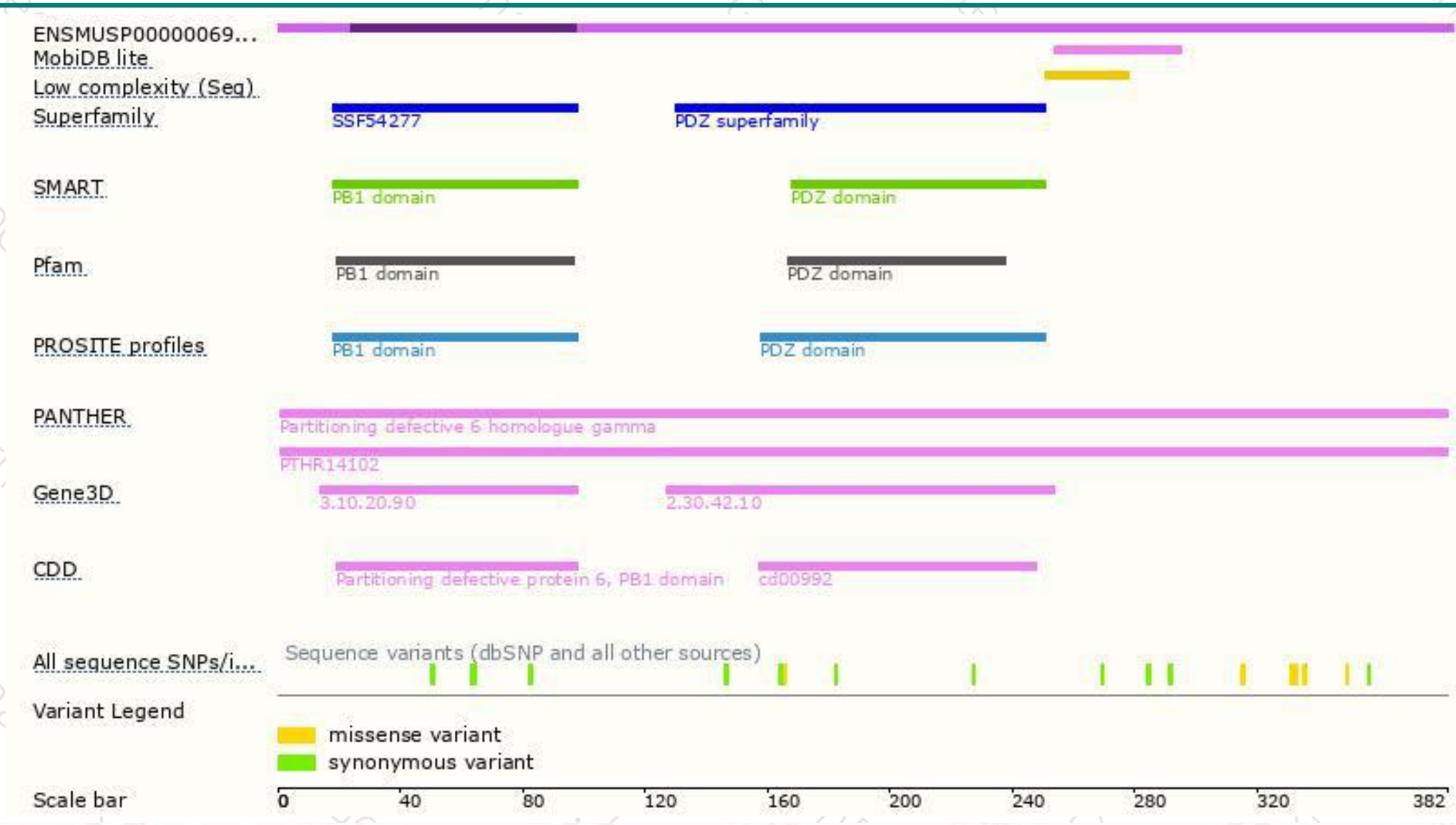
The strategy is based on the design of *Pard6g-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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