

Plekha8 Cas9-KO Strategy

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Project Overview

Project Name

Plekha8

Project type

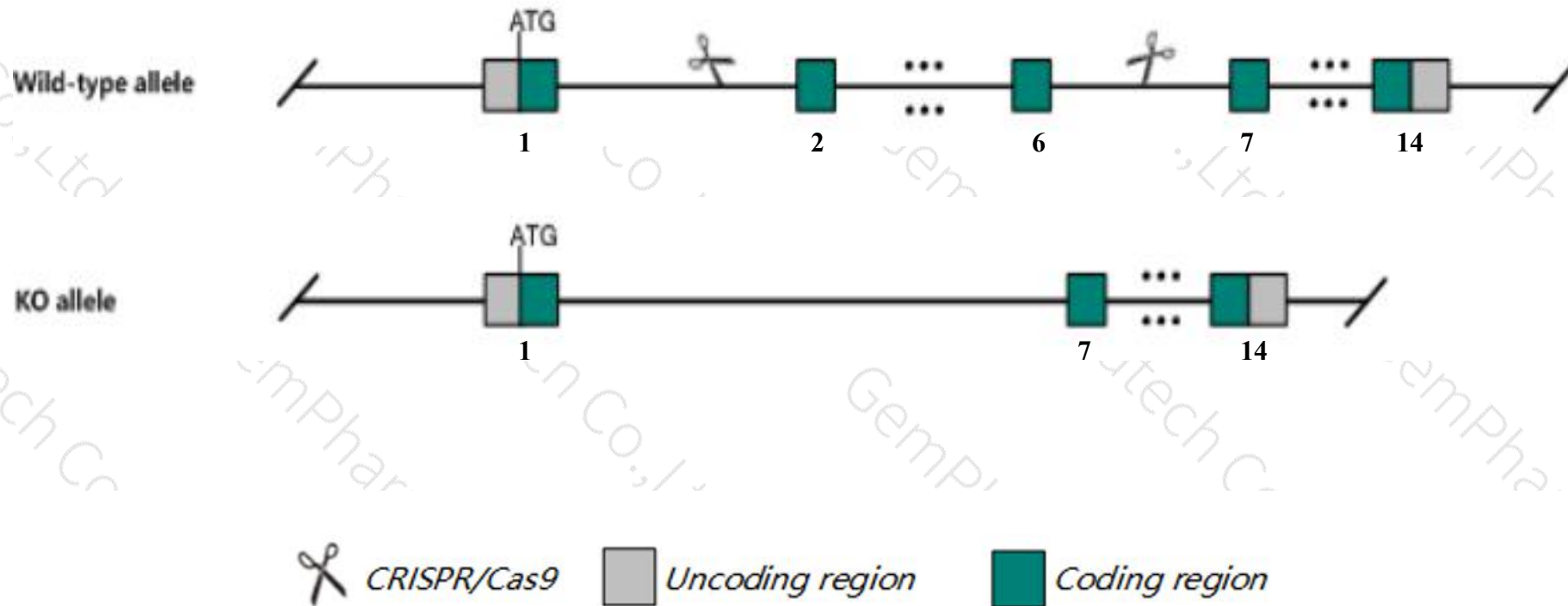
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Plekha8* gene. The schematic diagram is as follows:



Technical routes

- The *Plekha8* gene has 2 transcripts. According to the structure of *Plekha8* gene, exon2-exon6 of *Plekha8*-202(ENSMUST00000119706.7) transcript is recommended as the knockout region. The region contains 598bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Plekha8* gene. The brief process is as follows: CRISPR/Cas9 system were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

- The *Plekha8* gene is located on the Chr6. 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)

Plekha8 pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8 [Mus musculus (house mouse)]

Gene ID: 231999, updated on 13-Mar-2020

Summary



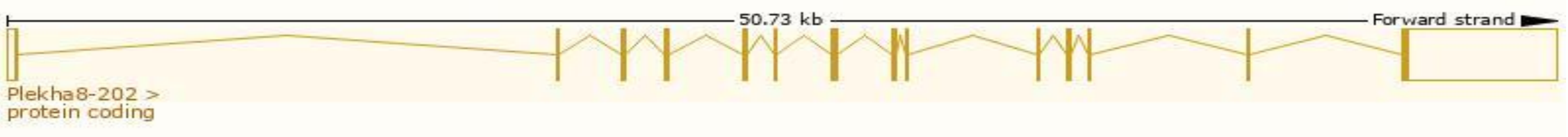
Official Symbol	Plekha8 provided by MGI
Official Full Name	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8 provided by MGI
Primary source	MGI:MGI:2681164
See related	Ensembl:ENSMUSG00000005225
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	6330400G01, AA517676, BC052360, Fapp-2, Fapp2
Expression	Ubiquitous expression in CNS E11.5 (RPKM 6.6), colon adult (RPKM 5.8) and 27 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

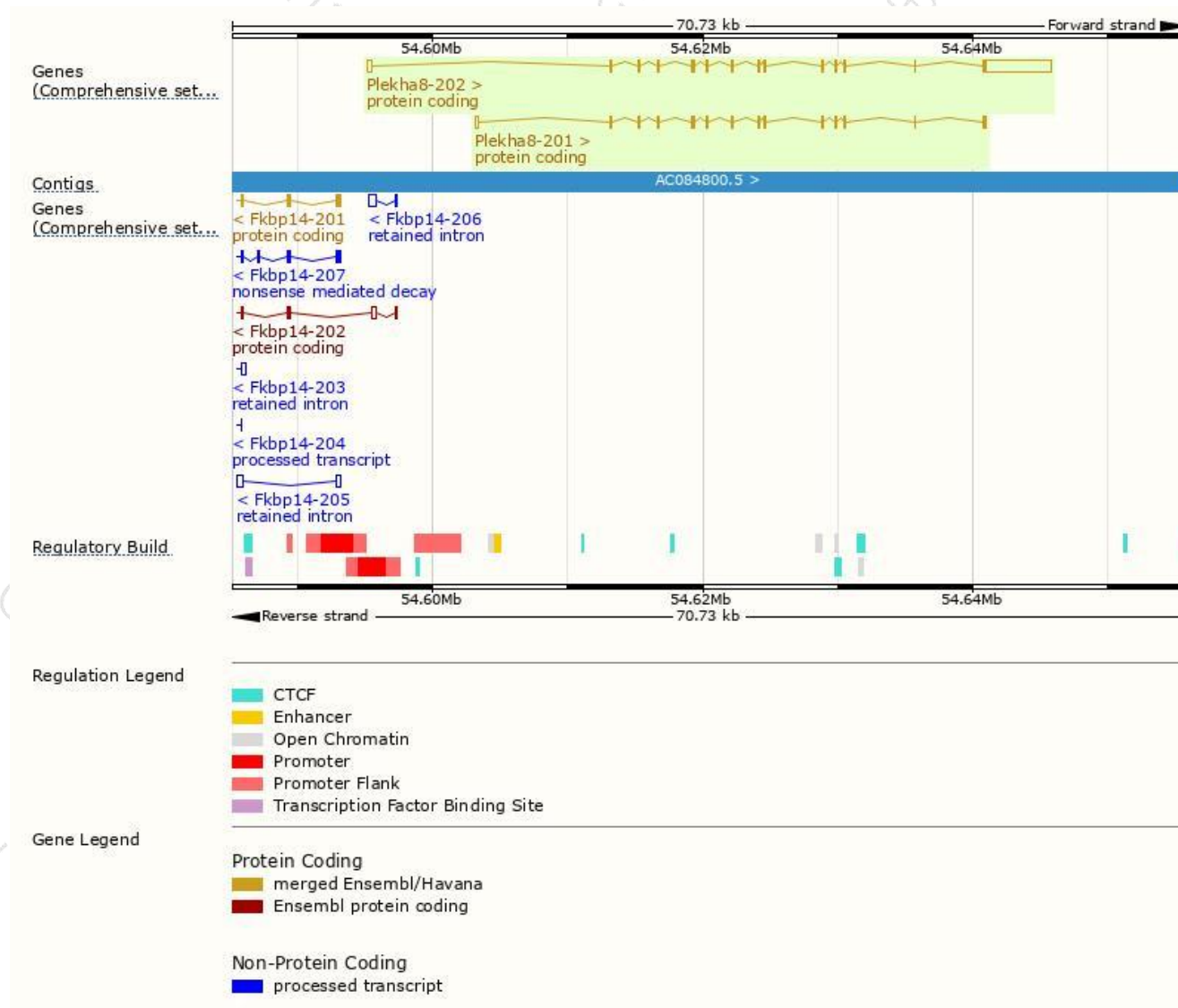
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Plekha8-202	ENSMUST00000119706.7	6739	519aa	Protein coding	CCDS51781	Q80W71	TSL:2 GENCODE basic APPRIS P1
Plekha8-201	ENSMUST00000101385.2	1838	474aa	Protein coding	CCDS39491	Q80W71	TSL:1 GENCODE basic

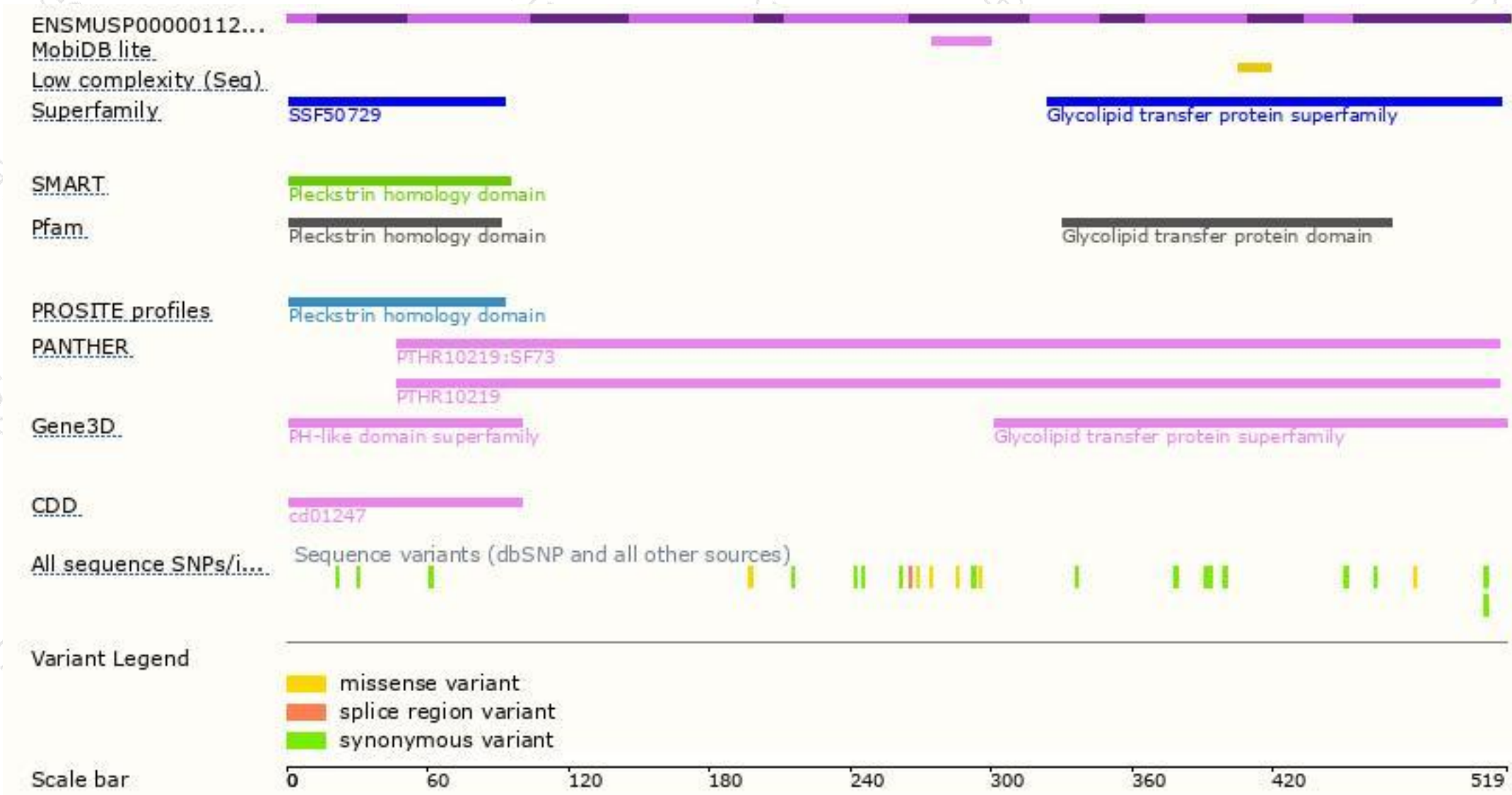
The strategy is based on the design of *Plekha8-202* transcript,the transcription is shown below:



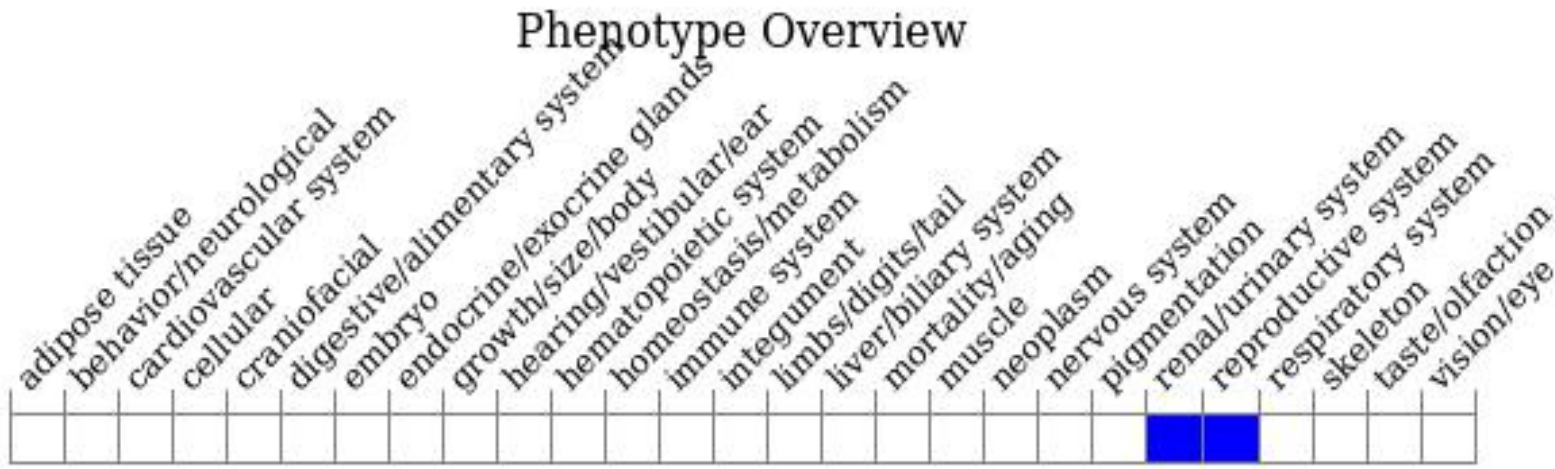
Genomic location distribution



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/>).

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

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