

Klhl34 Cas9-KO Strategy

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Project Overview

Project Name

Klhl34

Project type

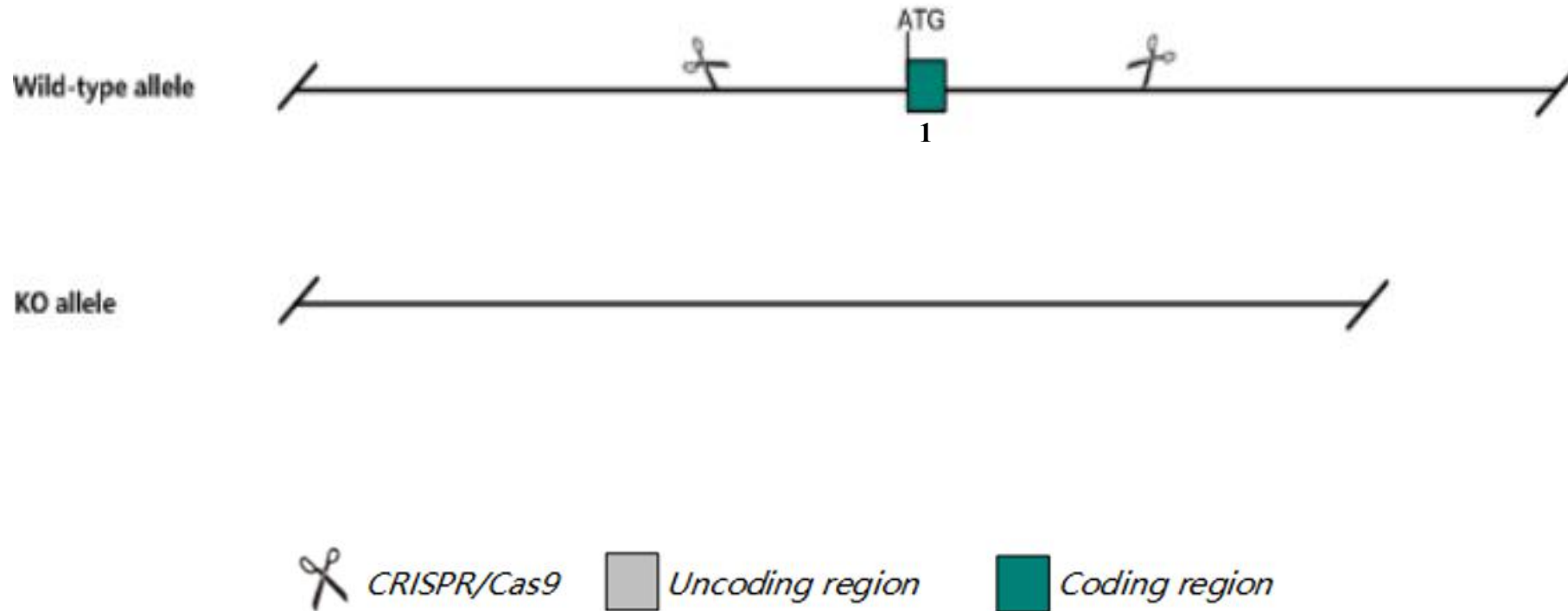
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Klhl34* gene. The schematic diagram is as follows:



The *Klhl34* gene has 1 transcript. According to the structure of *Klhl34* gene, exon1 of *Klhl34-201*(ENSMUST00000087157.4) 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 *Klhl34* 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 *Klhl34* gene is located on the ChrX. 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.

Klhl34 kelch-like 34 [Mus musculus (house mouse)]

Gene ID: 245683, updated on 13-Mar-2020

Summary

Official Symbol	Klhl34 provided by MGI
Official Full Name	kelch-like 34 provided by MGI
Primary source	MGI:MGI:2685234
See related	Ensembl:ENSMUSG00000047485
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	Gm388
Orthologs	human all

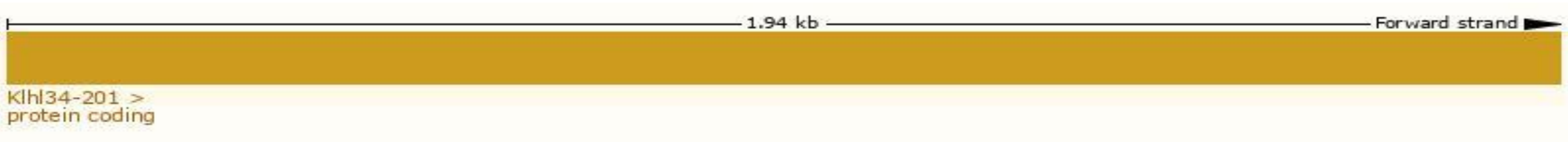
Transcript information Ensembl



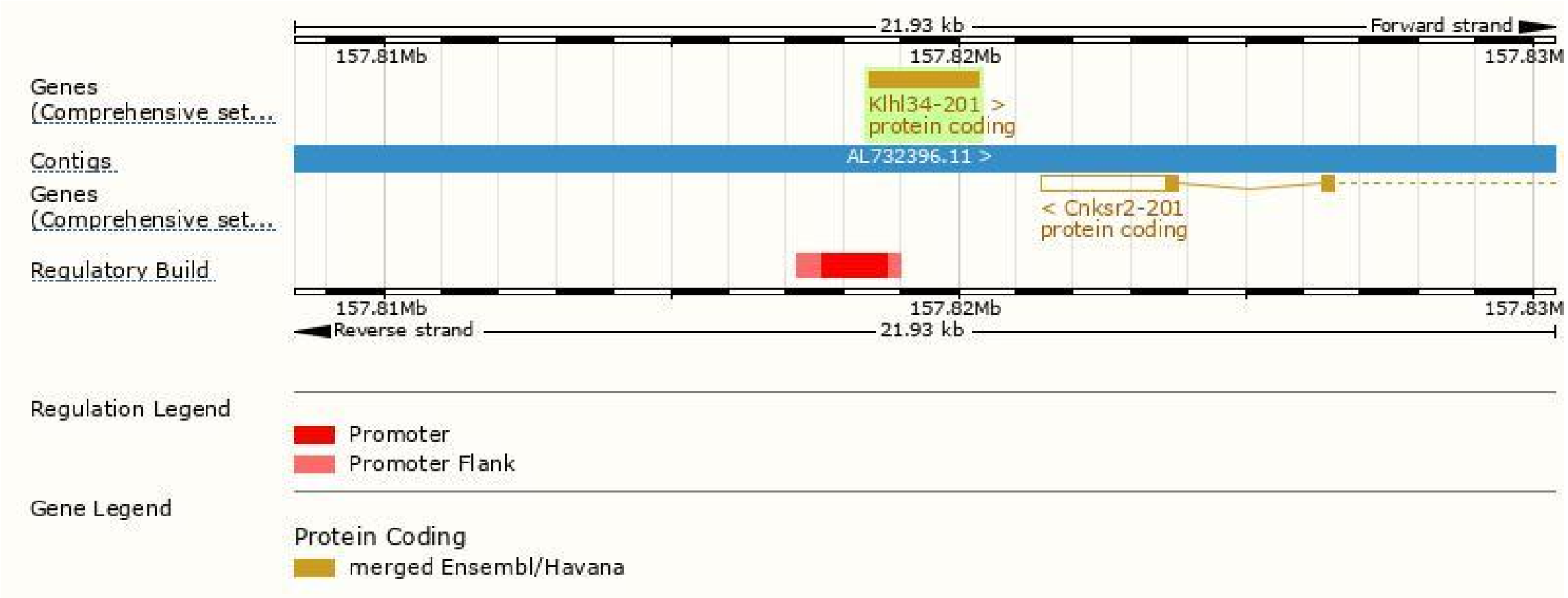
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Klhl34-201	ENSMUST00000087157.4	1935	644aa	Protein coding	CCDS41191	A2AI76	TSL:NA GENCODE basic APPRIS P1

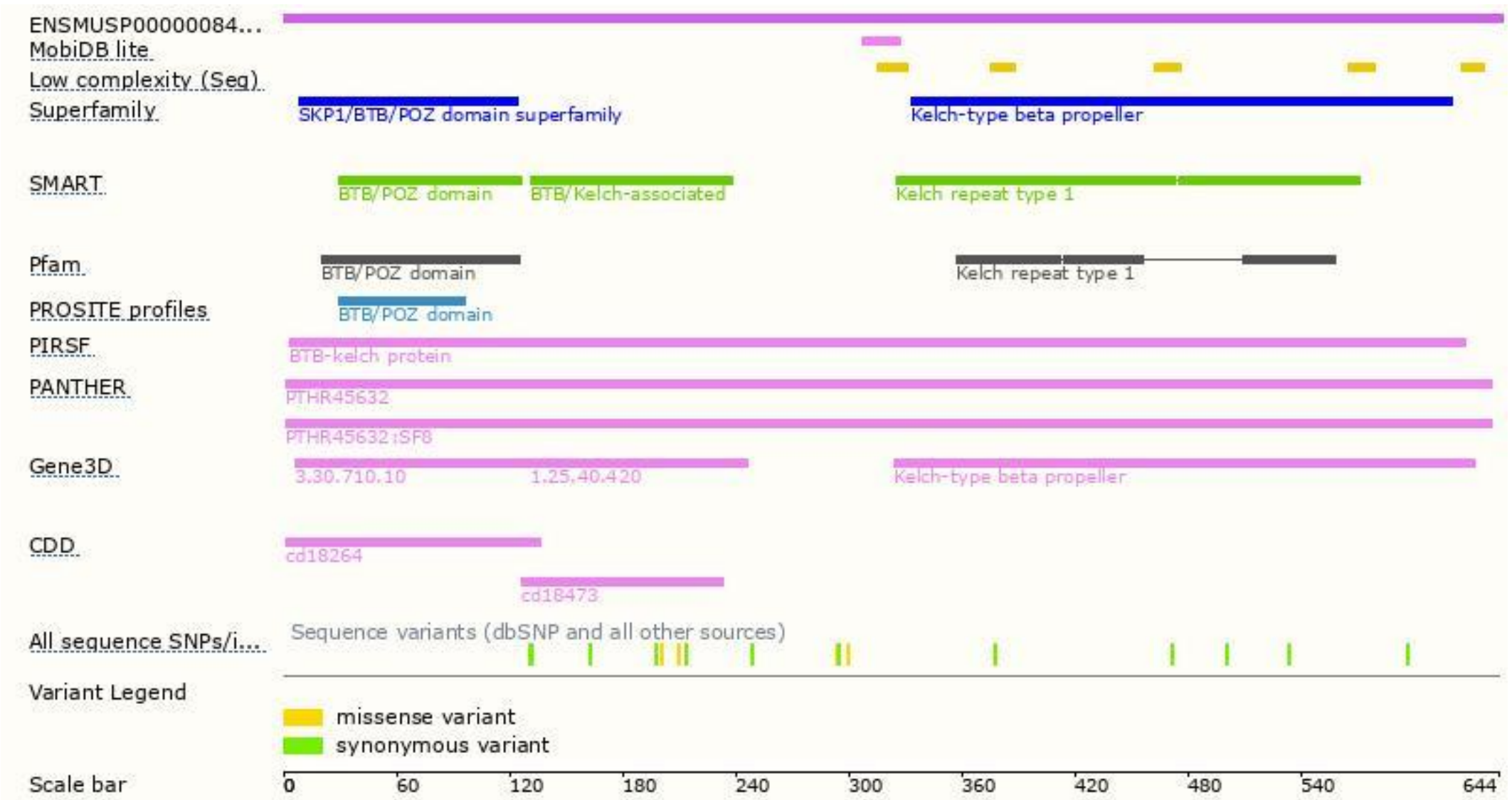
The strategy is based on the design of *Klhl34-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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