

Dnaaf1 Cas9-KO Strategy

Designer: Xiaojing Li

Reviewer: Jia Yu

Design Date: 2020-4-20

Project Overview

Project Name

Dnaaf1

Project type

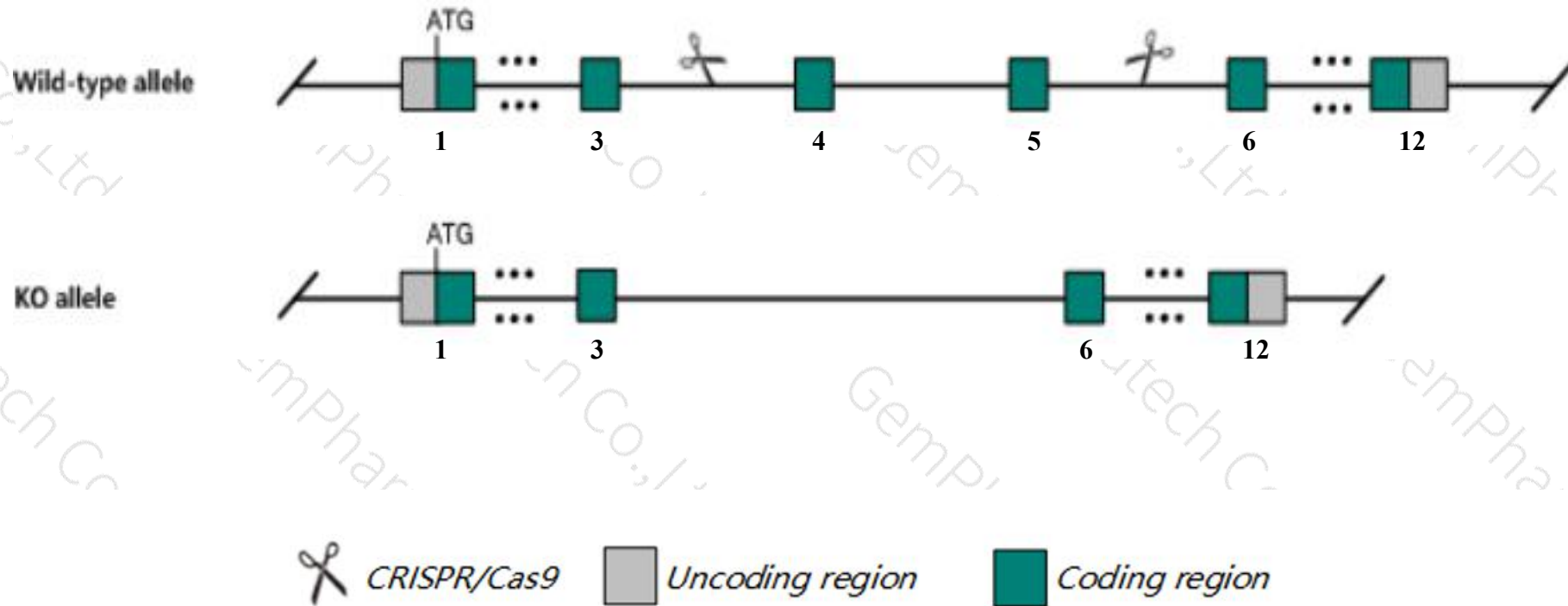
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dnaaf1* gene. The schematic diagram is as follows:



- The *Dnaaf1* gene has 1 transcript. According to the structure of *Dnaaf1* gene, exon4-exon5 of *Dnaaf1*-201 (ENSMUST00000093100.2) transcript is recommended as the knockout region. The region contains 389bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dnaaf1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, mice homozygous for an enu-induced mutation exhibit partial postnatal lethality, domed cranium, enlarged lateral ventricles, abnormal hippocampus morphology, and thin cerebral cortex.
- The *Dnaaf1* gene is located on the Chr8. 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)

Dnaaf1 dynein, axonemal assembly factor 1 [Mus musculus (house mouse)]

Gene ID: 68270, updated on 13-Mar-2020

Summary



Official Symbol	Dnaaf1 provided by MGI
Official Full Name	dynein, axonemal assembly factor 1 provided by MGI
Primary source	MGI:MGI:1915520
See related	Ensembl:ENSMUSG00000031831
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	4930457P18Rik, Lrrc50, m4Bei
Expression	Biased expression in testis adult (RPKM 269.9) and duodenum adult (RPKM 6.3) See more
Orthologs	human all

Transcript information (Ensembl)

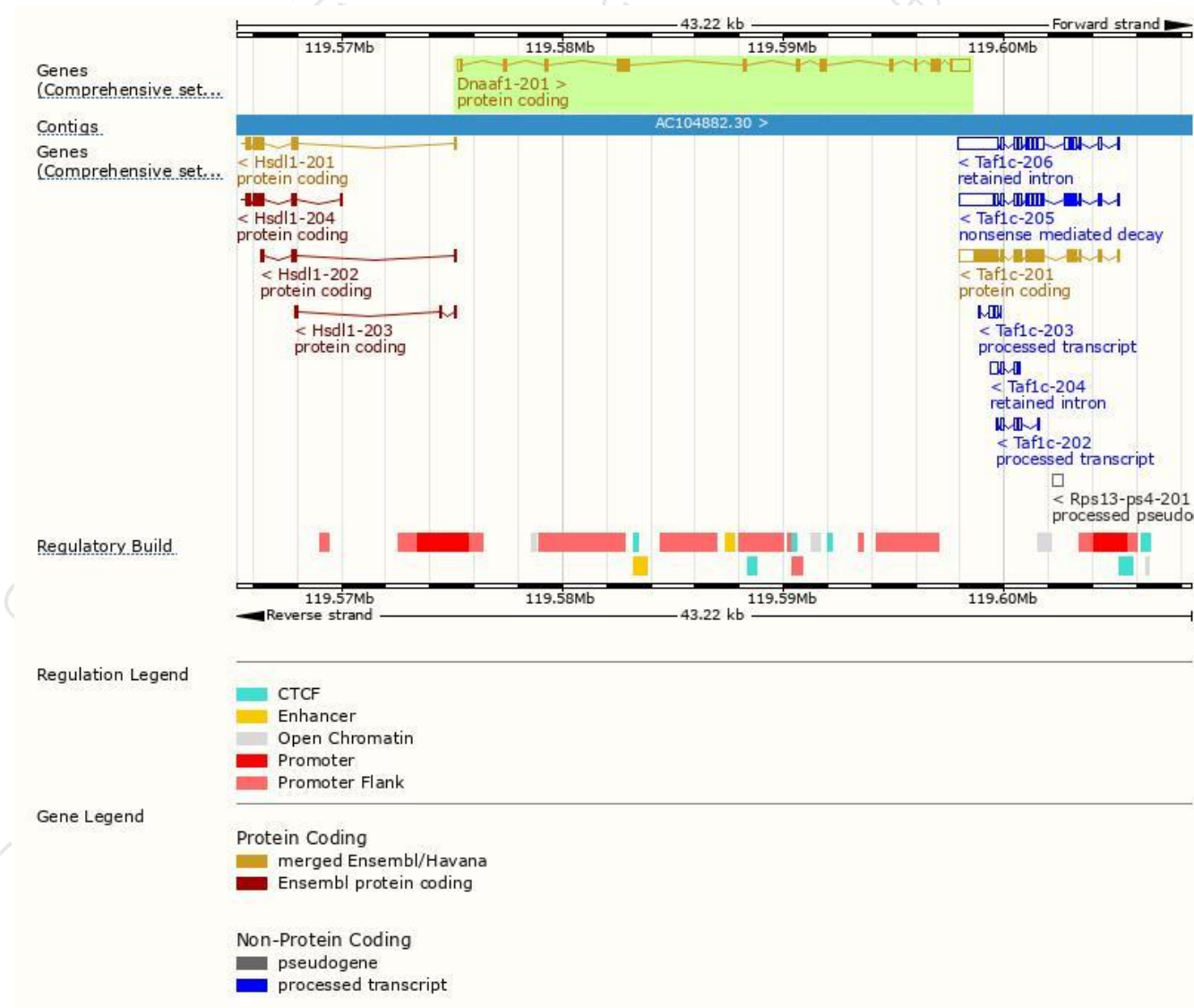
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dnaaf1-201	ENSMUST00000093100.2	2821	634aa	Protein coding	CCDS22709	Q9D2H9	TSL:1 GENCODE basic APPRIS P1

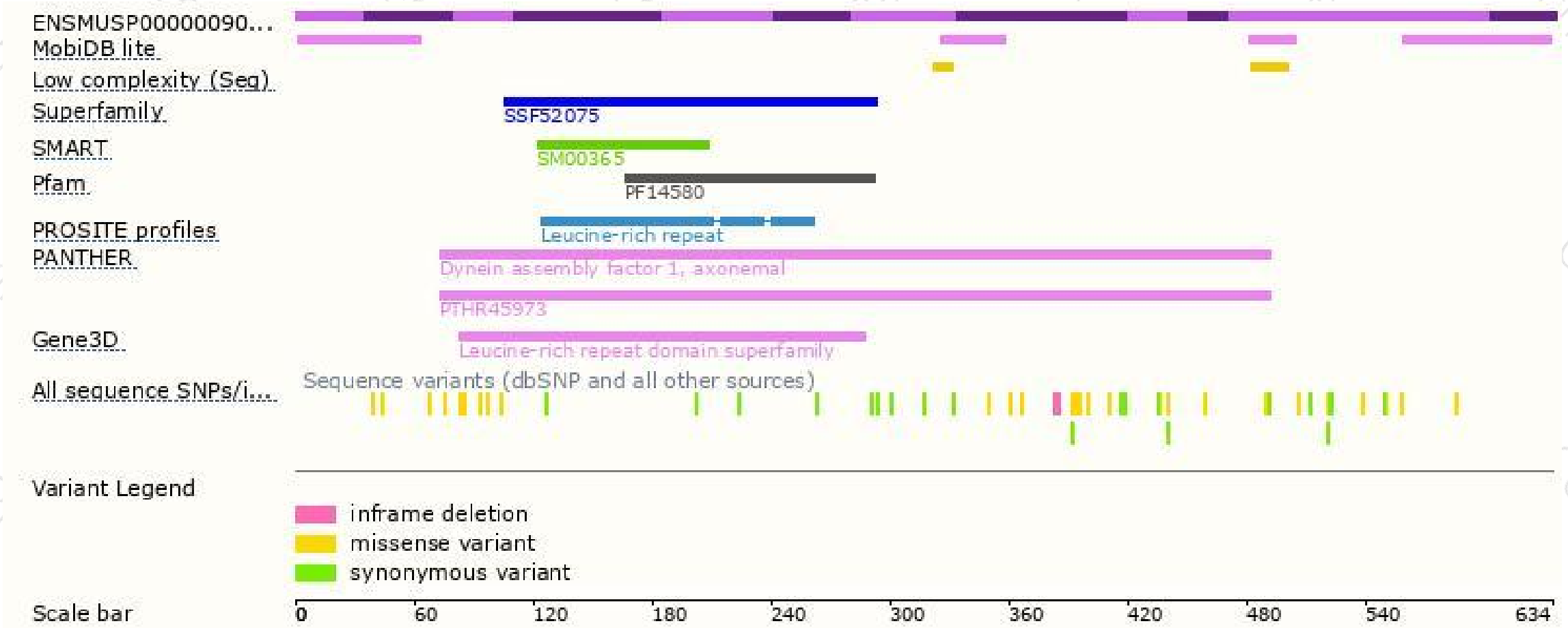
The strategy is based on the design of *Dnaaf1-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

Tel: 400-9660890

