

Dok5 Cas9-KO Strategy

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Project Overview

Project Name

Dok5

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dok5* gene. The schematic diagram is as follows:



- The *Dok5* gene has 1 transcript. According to the structure of *Dok5* gene, exon3-exon4 of *Dok5-201* (ENSMUST00000029075.4) transcript is recommended as the knockout region. The region contains 235bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dok5* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Dok5* gene is located on the Chr2. 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)

Dok5 docking protein 5 [Mus musculus (house mouse)]

Gene ID: 76829, updated on 13-Mar-2020

Summary



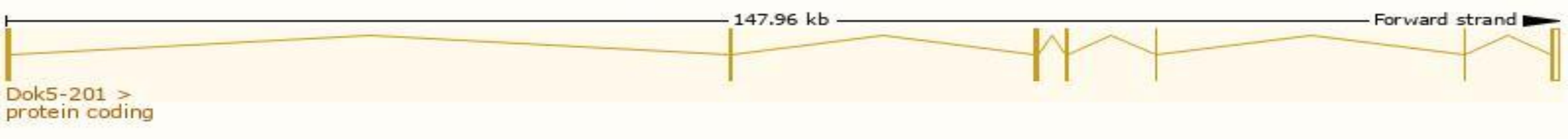
Official Symbol	Dok5 provided by MGI
Official Full Name	docking protein 5 provided by MGI
Primary source	MGI:MGI:1924079
See related	Ensembl:ENSMUSG000000027560
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2700055C10Rik
Expression	Biased expression in CNS E18 (RPKM 19.0), CNS E14 (RPKM 9.0) and 7 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

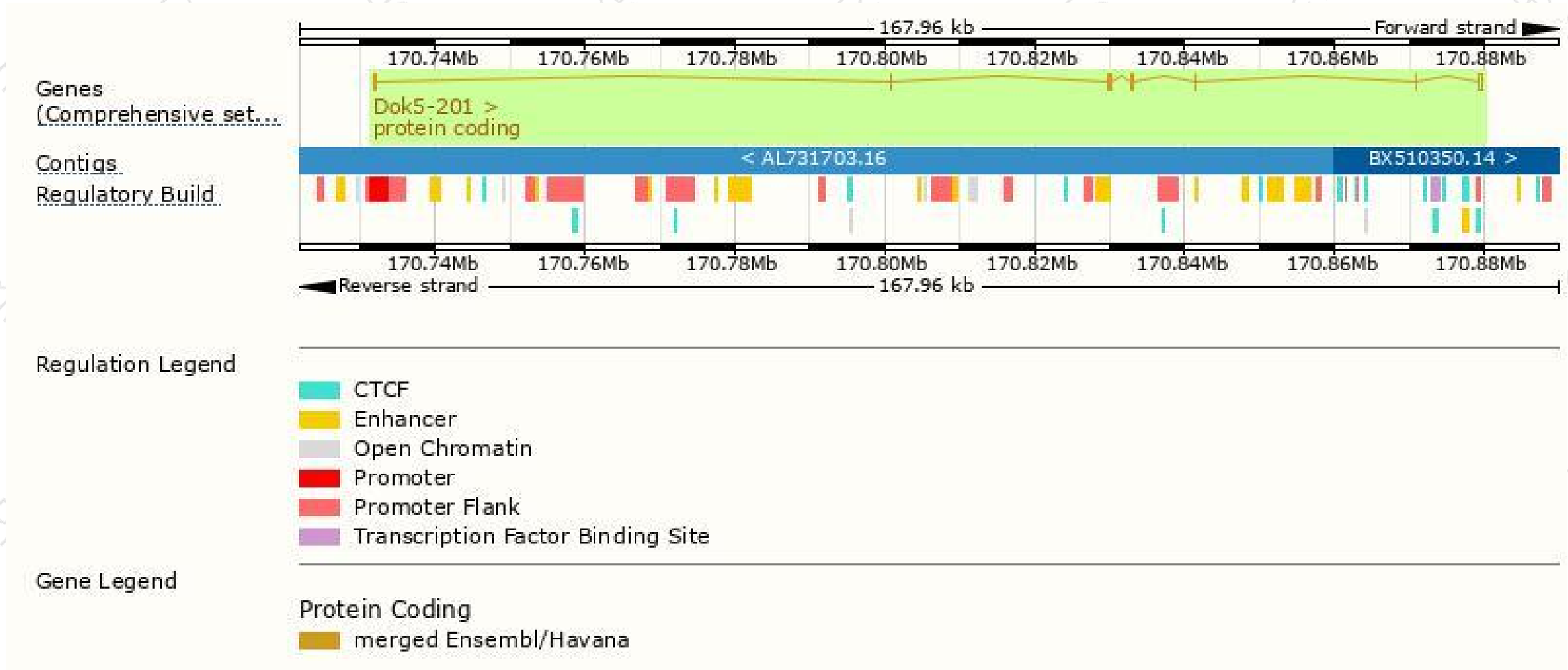
The gene has 1 transcript, and the transcript is shown below:

Name ▲	Transcript ID ▲	bp ▲	Protein ▲	Biotype ▲	CCDS ▲	UniProt ▲	Flags ▲
Dok5-201	ENSMUST00000029075.4	1768	306aa	Protein coding	CCDS17125	Q91ZM9	TSL:1 Gencode basic APPRIS P1

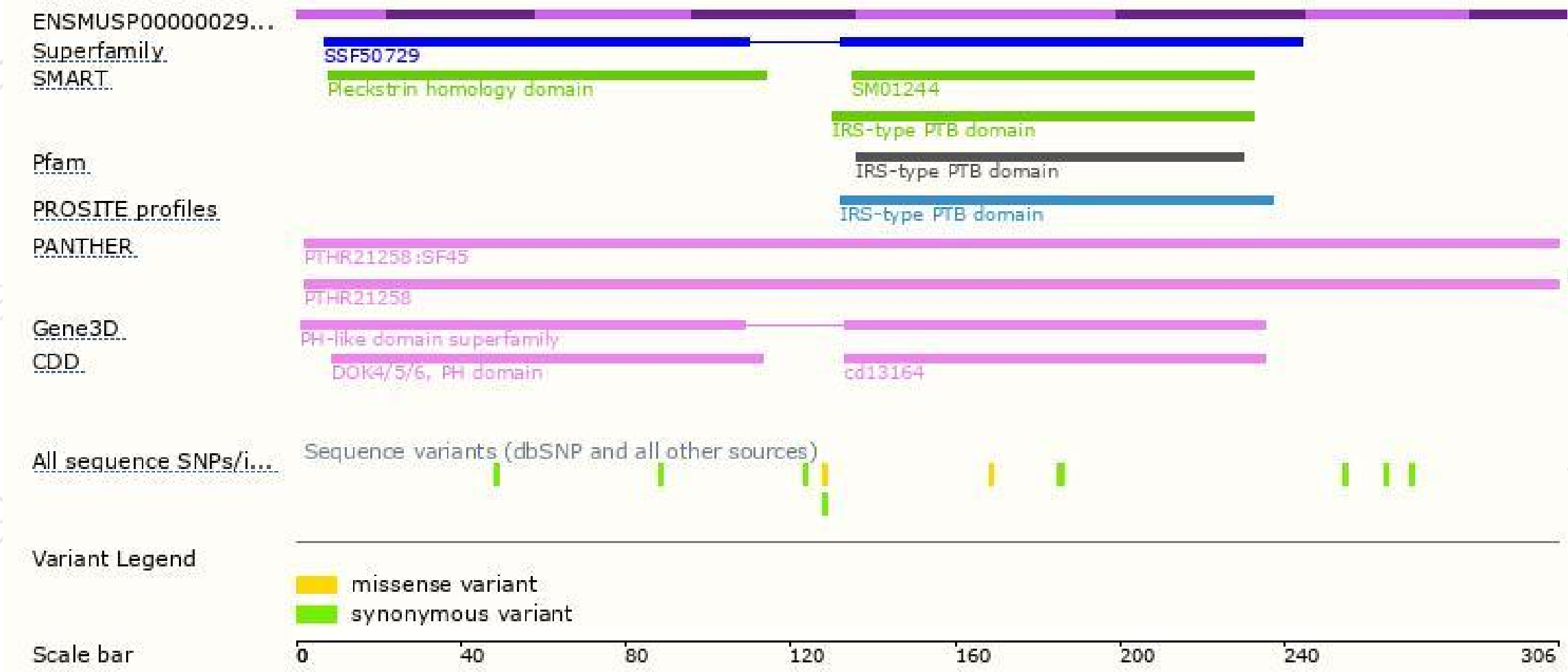
The strategy is based on the design of *Dok5-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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