

# Dok5 Cas9-CKO Strategy

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# **Project Overview**



**Project Name** 

Dok5

**Project type** 

Cas9-CKO

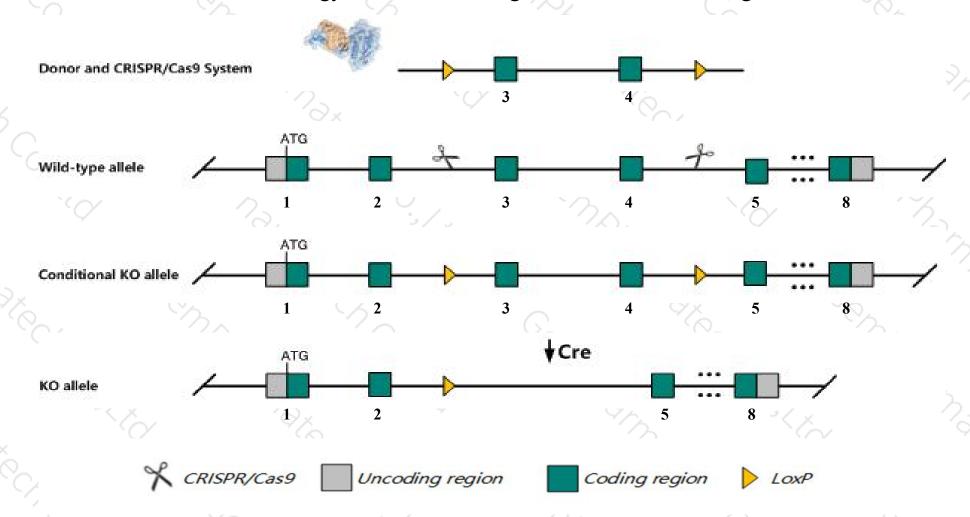
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



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



### Technical routes



- ➤ 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 and Donor 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 flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

### **Notice**



- > 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological 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:ENSMUSG00000027560

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 tissuesSee more

Orthologs <u>human</u> all

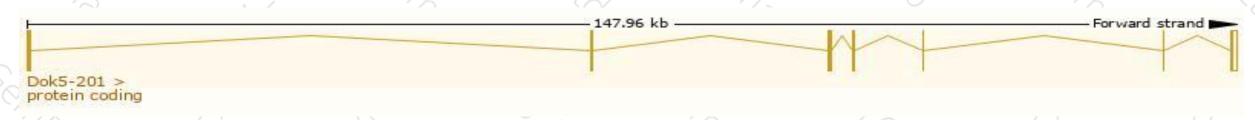
# Transcript information (Ensembl)



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

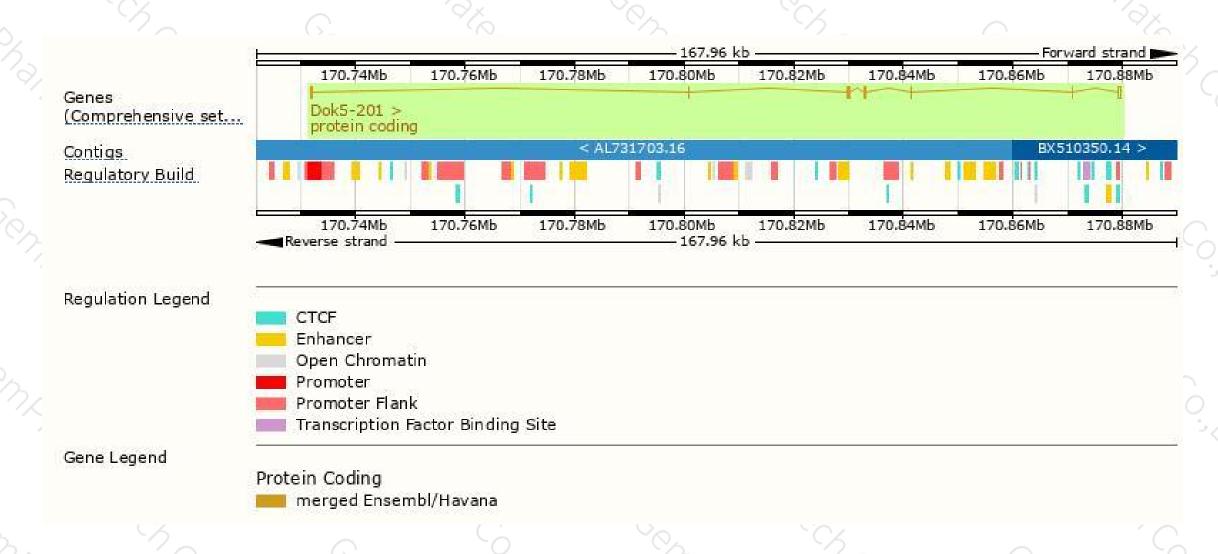
Name A	Transcript ID	bp 🌲	Protein #	Biotype 🍦	CCDS 🍦	UniProt #	Flags		
Dok5-201	ENSMUST00000029075.4	1768	<u>306aa</u>	Protein coding	CCDS17125₽	Q91ZM9₽	TSL:1	GENCODE basic	APPRIS P1

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



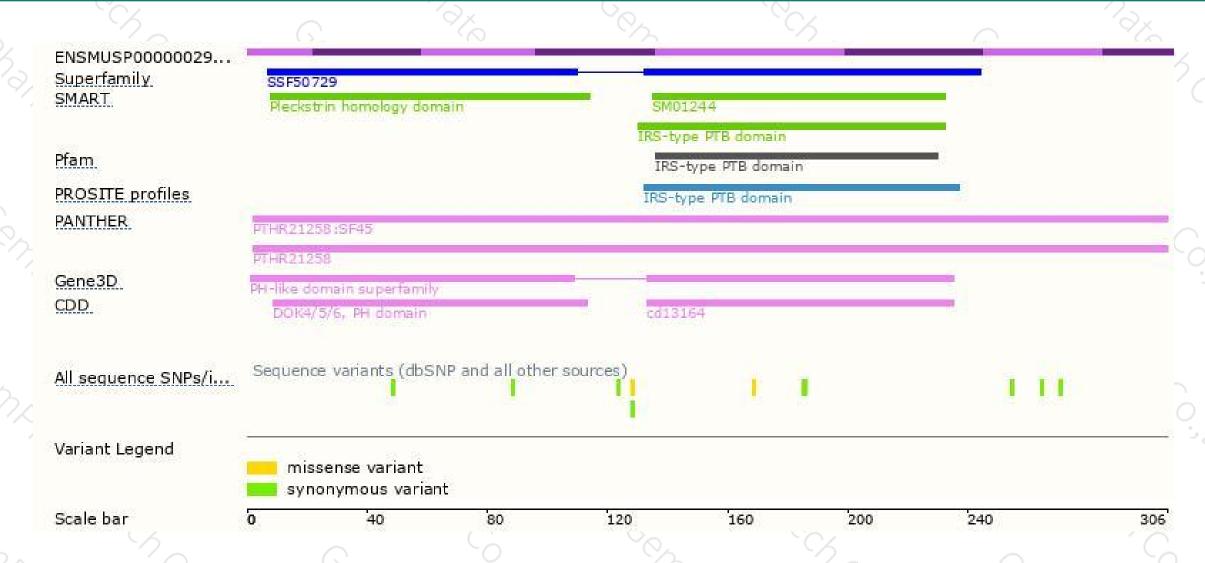
### Genomic location distribution





### Protein domain







If you have any questions, you are welcome to inquire. Tel: 400-9660890





