

Tab3 Cas9-CKO Strategy

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Project Overview

Project Name

Tab3

Project type

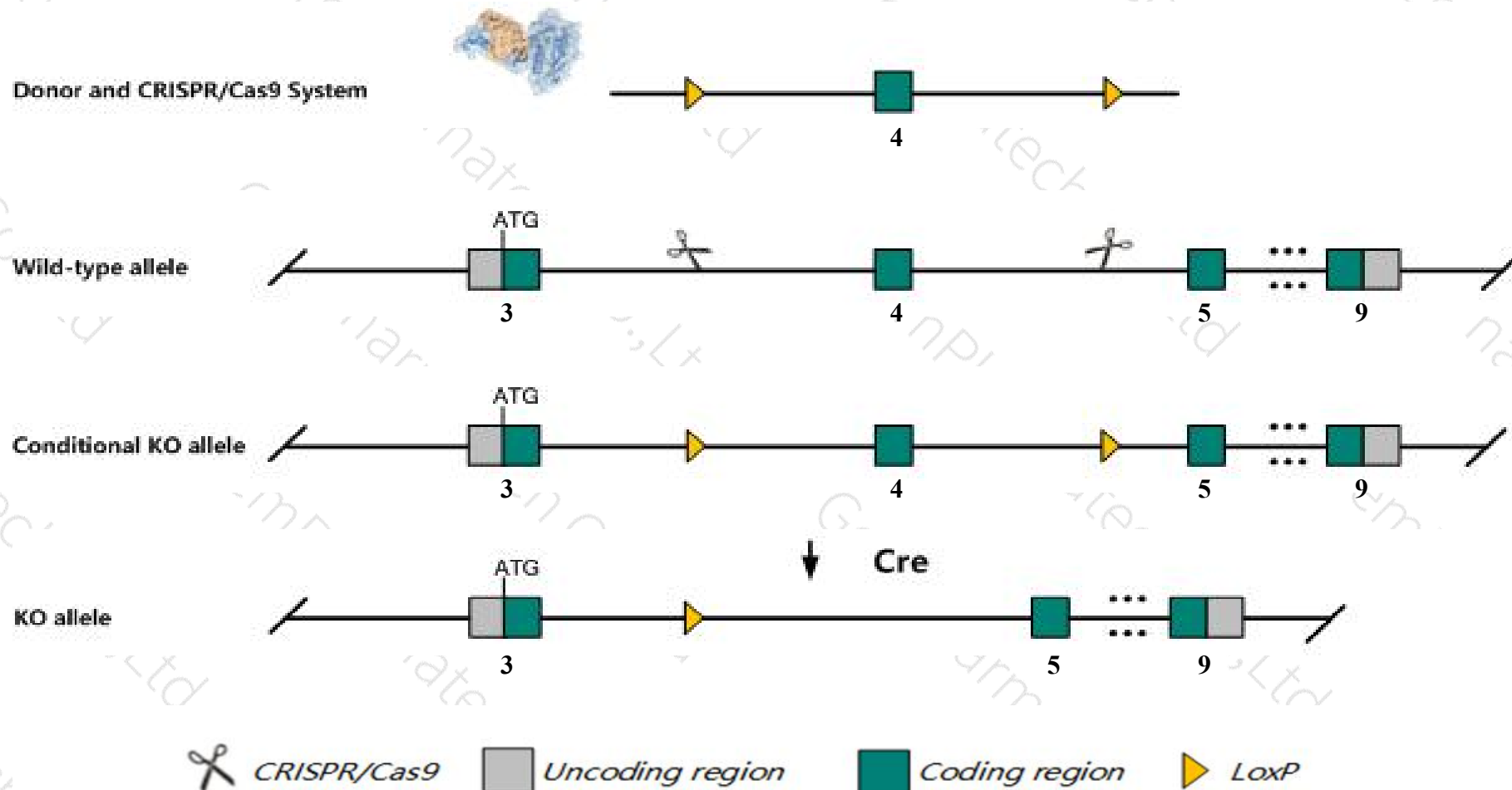
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Tab3* gene. The schematic diagram is as follows:



- The *Tab3* gene has 4 transcripts. According to the structure of *Tab3* gene, exon4 of *Tab3-201* (ENSMUST00000048250.9) transcript is recommended as the knockout region. The region contains 1459bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Tab3* 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.

- According to the existing MGI data, Male chimeras hemizygous for a gene trapped allele appear normal at E9.5.
- The *Tab3* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Tab3 TGF-beta activated kinase 1/MAP3K7 binding protein 3 [*Mus musculus* (house mouse)]

Gene ID: 66724, updated on 24-Oct-2019

Summary

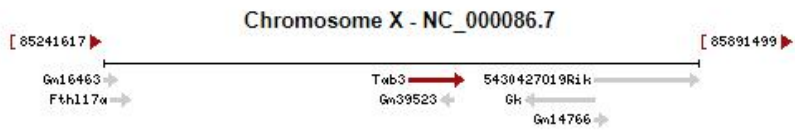
- Official Symbol** Tab3 provided by [MGI](#)
- Official Full Name** TGF-beta activated kinase 1/MAP3K7 binding protein 3 provided by [MGI](#)
- Primary source** [MGI:MGI:1913974](#)
- See related** [Ensembl:ENSMUSG00000035476](#)
- Gene type** protein coding
- RefSeq status** VALIDATED
- Organism** [Mus musculus](#)
- Lineage** Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
- Also known as** Map3k7ip3; mKIAA4135; 4921526G09Rik
- Expression** Ubiquitous expression in liver E14 (RPKM 4.3), placenta adult (RPKM 4.2) and 27 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: X; X C1 See Tab3 in [Genome Data Viewer](#)

Exon count: 11

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	X	NC_000086.7 (85573854..85634469)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	X	NC_000086.6 (82819361..82879808)



Transcript information (Ensembl)

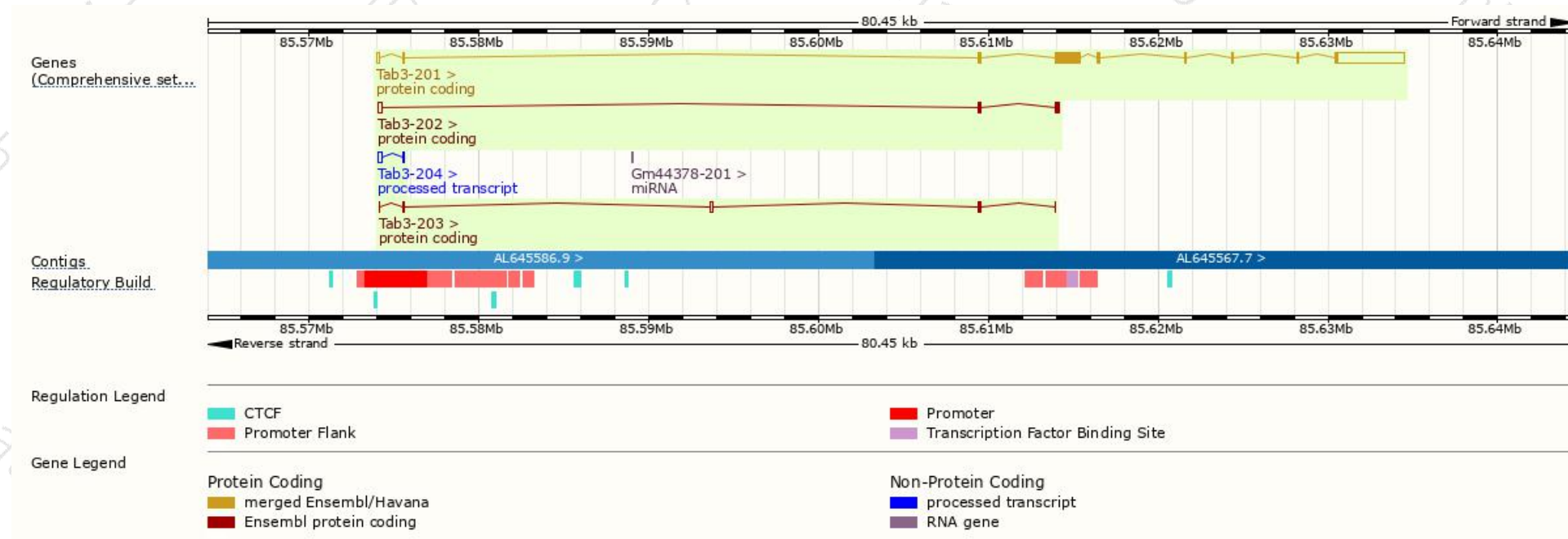
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tab3-201	ENSMUST00000048250.9	6364	716aa	Protein coding	CCDS41049	B1ASY9 Q571K4	TSL:1 GENCODE basic APPRIS P1
Tab3-202	ENSMUST00000137438.1	707	108aa	Protein coding	-	F7AT87	CDS 3' incomplete TSL:2
Tab3-203	ENSMUST00000146063.1	385	38aa	Protein coding	-	F6XIV6	CDS 3' incomplete TSL:5
Tab3-204	ENSMUST00000152880.1	360	No protein	Processed transcript	-	-	TSL:3

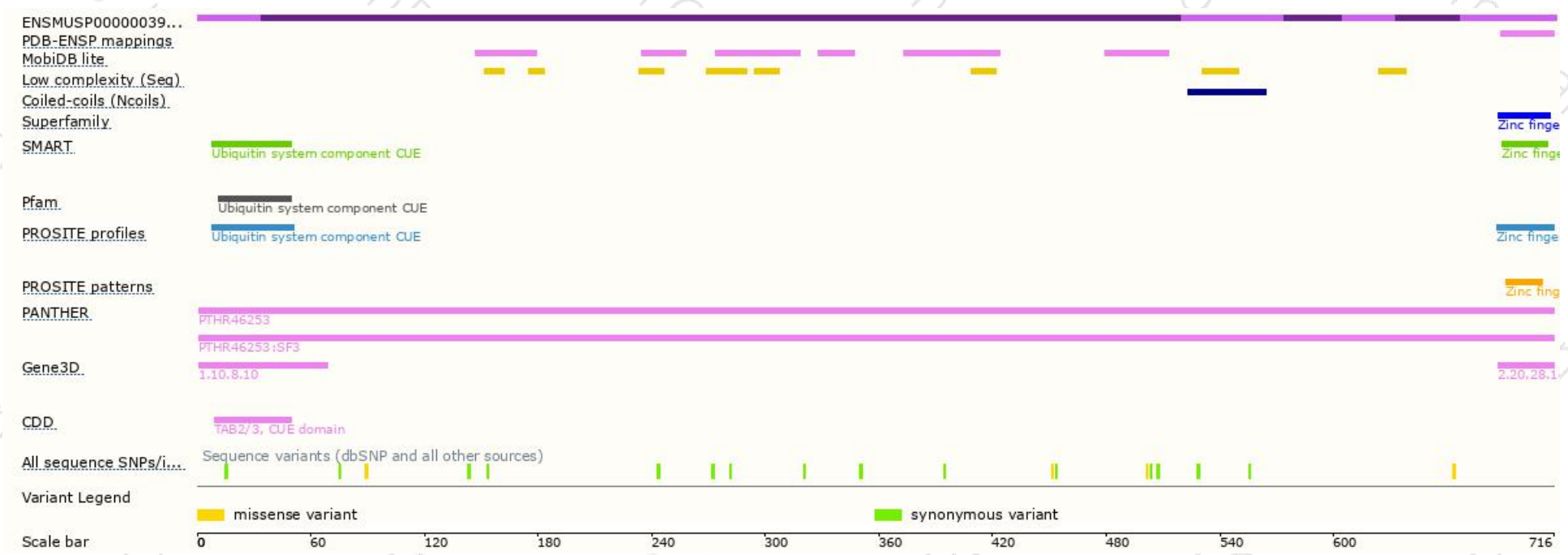
The strategy is based on the design of *Tab3-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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