

# ***Trim69 Cas9-CKO Strategy***

**Designer:** Huan Wang  
**Reviewer:** Huan Fan  
**Design Date:** 2019-11-19

# Project Overview

**Project Name**

***Trim69***

**Project type**

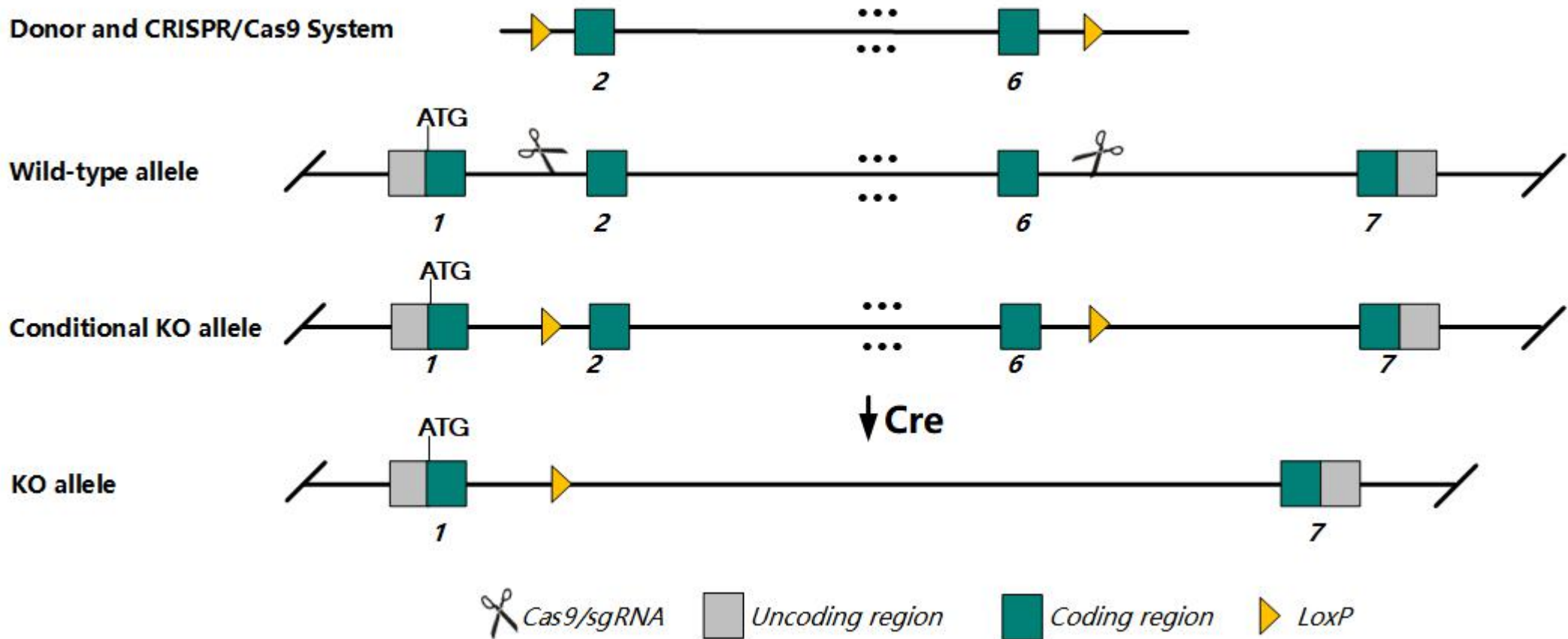
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Trim69* gene. The schematic diagram is as follows:



# Technical routes

- The *Trim69* gene has 2 transcript. According to the structure of *Trim69* gene, exon2-6 of *Trim69*-201( ENSMUST00000036089.7) transcript is recommended as the knockout region. The region contains 958bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Trim69* gene. The brief process is as follows: gRNA was transcribed in vitro, donor was constructed. Cas9, gRNA 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 or cell types.

- The KO region contains functional region of the *Patl2* gene. Knockout the region may affect the function of *Patl2* gene.
- The *Trim69* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.



# Gene information ( NCBI )

## Trim69 tripartite motif-containing 69 [ *Mus musculus* (house mouse) ]

Gene ID: 70928, updated on 12-Aug-2019

### Summary

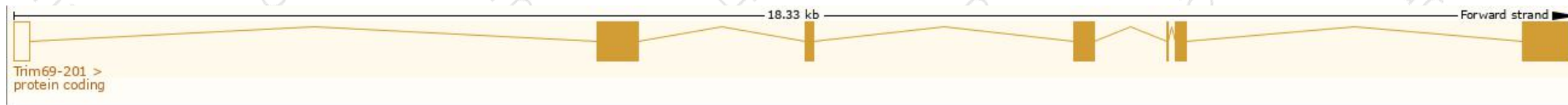
<b>Official Symbol</b>	Trim69 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	tripartite motif-containing 69 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1918178</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000033368</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	Trif, Rnf36, Trimless; 4921519C19Rik
<b>Expression</b>	Restricted expression toward testis adult (RPKM 46.9) <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information ( Ensembl )

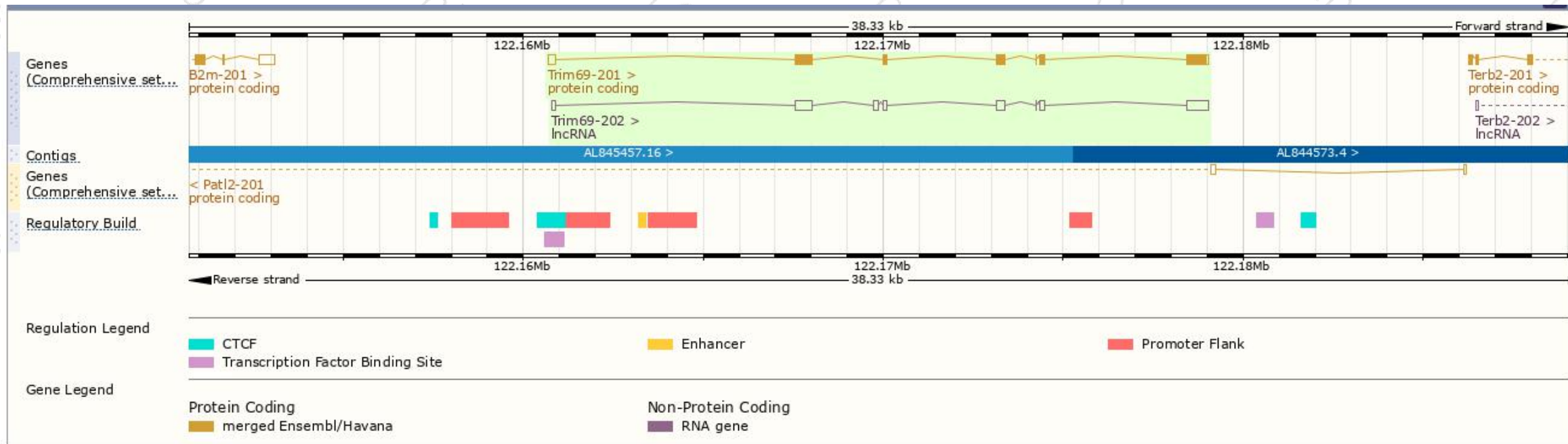
The gene has 2 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Trim69-201	<a href="#">ENSMUST0000036089.7</a>	1758	<a href="#">500aa</a>	Protein coding	<a href="#">CCDS16655</a>	<a href="#">Q80X56</a>	TSL:1 GENCODE basic APPRIS P1
Trim69-202	<a href="#">ENSMUST00000143088.1</a>	1783	No protein	lncRNA	-	-	TSL:5

The strategy is based on the design of *Trim69-201* transcript, The transcription is shown below



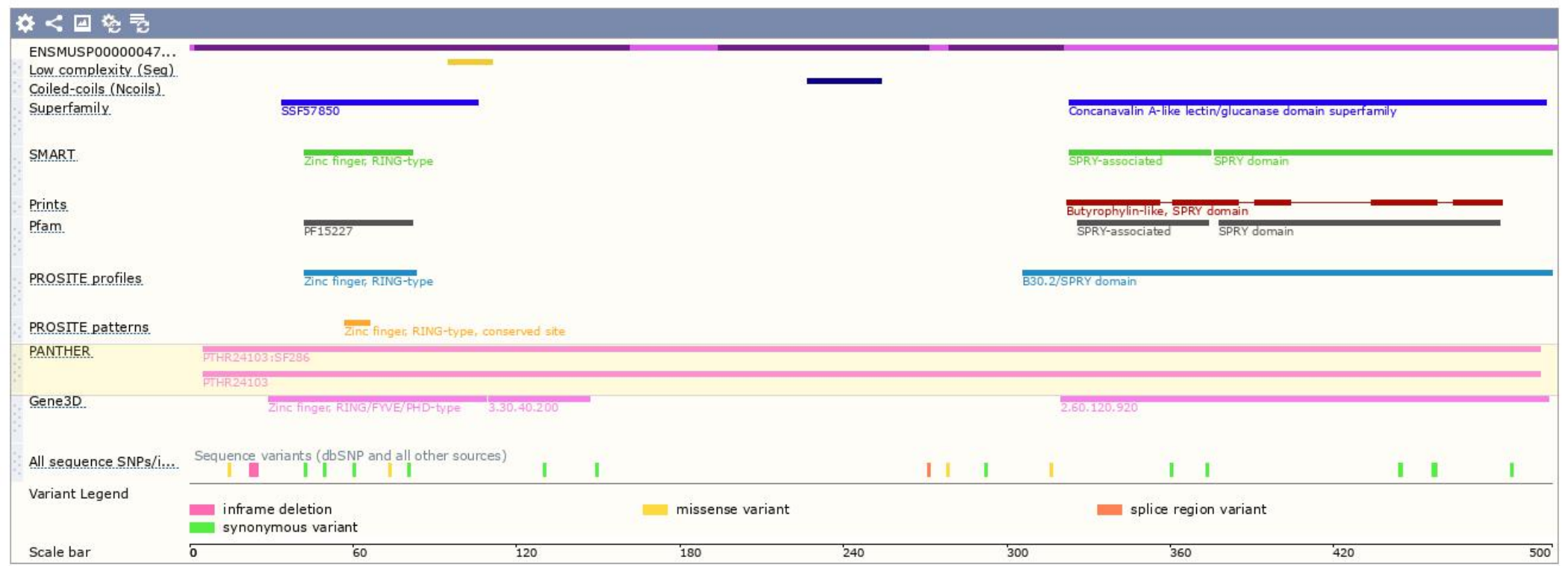
# Genomic location distribution





# Protein domain

Protein domains for ENSMUSP00000047627.7



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

