

Trem12 Cas9-CKO Strategy

Designer:

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Project Overview

Project Name

Trem12

Project type

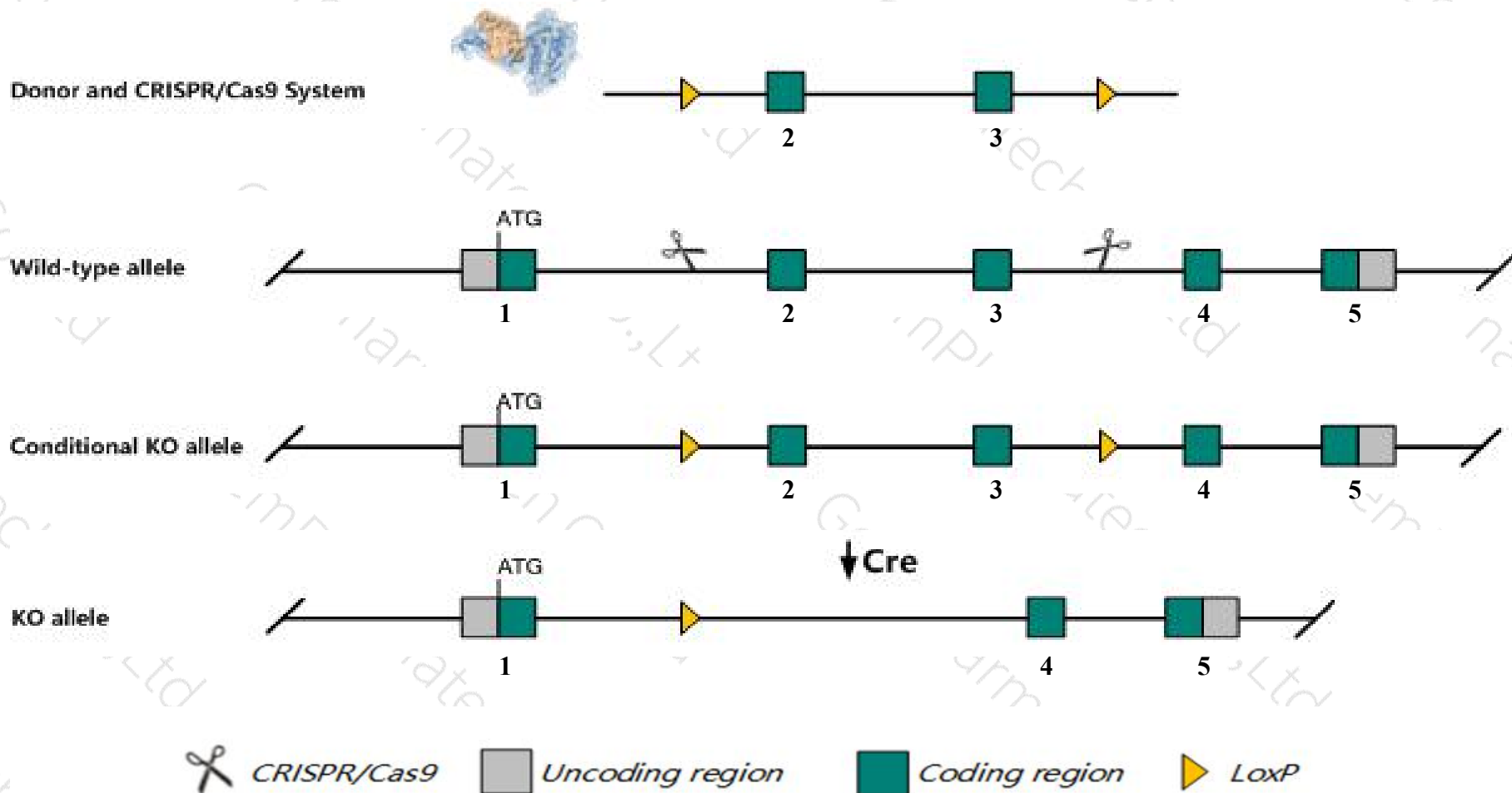
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Trem12* gene. The schematic diagram is as follows:



- The *Trem12* gene has 4 transcripts. According to the structure of *Trem12* gene, exon2-exon3 of *Trem12*-202 (ENSMUST00000233092.1) transcript is recommended as the knockout region. The region contains 715bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Trem12* 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 *Trem12* gene is located on the Chr17. 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)

Trem12 triggering receptor expressed on myeloid cells-like 2 [Mus musculus (house mouse)]

Gene ID: 328833, updated on 31-Jan-2019

Summary



Official Symbol	Trem12 provided by MGI
Official Full Name	triggering receptor expressed on myeloid cells-like 2 provided by MGI
Primary source	MGI:MGI:2147038
See related	Ensembl:ENSMUSG00000071068
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	AW049306, Gm750, Tlt2
Expression	Biased expression in spleen adult (RPKM 15.4), thymus adult (RPKM 10.4) and 5 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

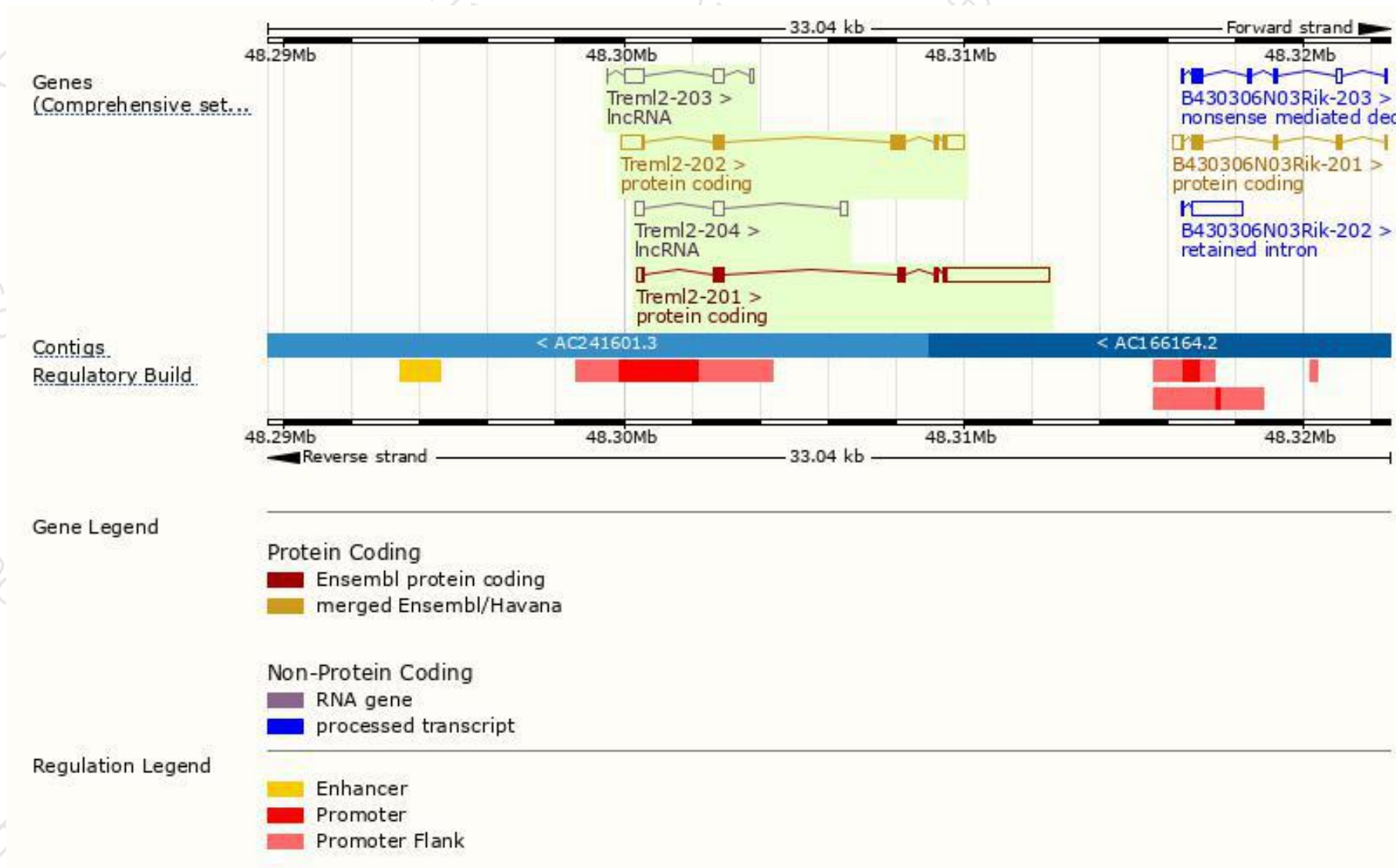
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Trem12-202	ENSMUST00000233092.1	2094	329aa	Protein coding	CCDS37646	Q2LA85	GENCODE basic APPRIS P2
Trem12-201	ENSMUST00000170941.2	3983	270aa	Protein coding	-	A0A3F2YNQ7	TSL:1 GENCODE basic APPRIS ALT2
Trem12-203	ENSMUST00000233496.1	984	No protein	lncRNA	-	-	
Trem12-204	ENSMUST00000233602.1	770	No protein	lncRNA	-	-	

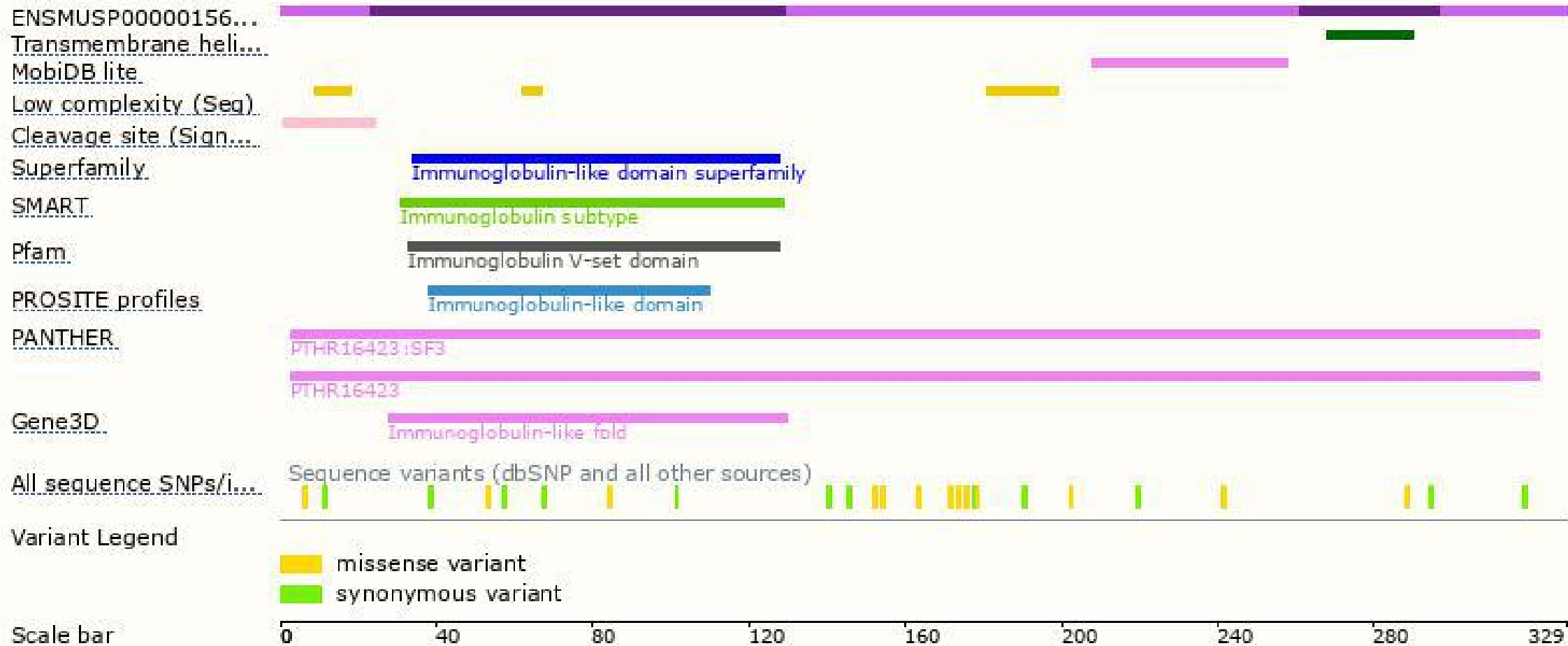
The strategy is based on the design of *Trem12-202* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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