

Mbl2 Cas9-CKO Strategy

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Overview

Target Gene Name

• Mbl2

Project Type

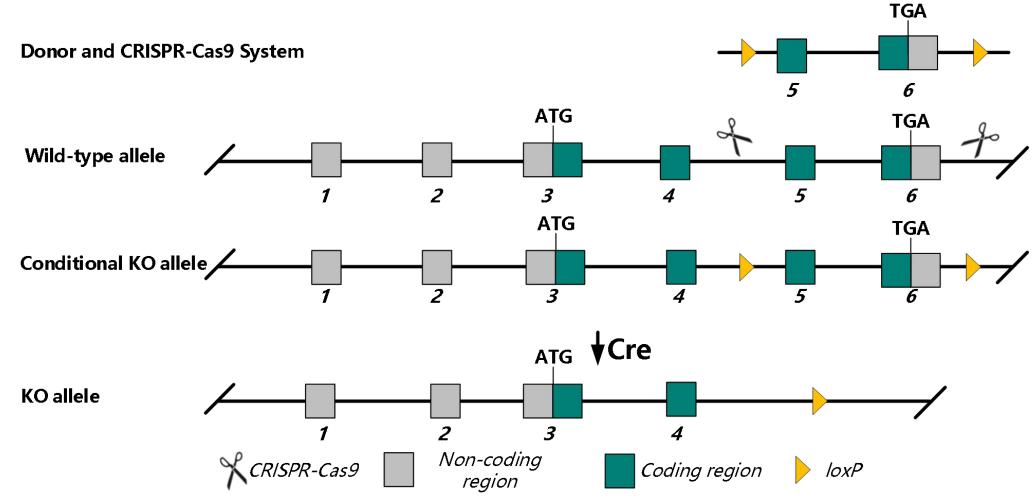
• Cas9-CKO

Genetic Background

• C57BL/6JGpt



Strain Strategy

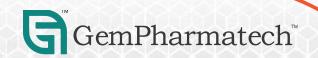


Schematic representation of CRISPR-Cas9 engineering used to edit the Mbl2 gene.



Technical Information

- The *Mbl2* gene has 1 transcript. According to the structure of *Mbl2* gene, exon 5-6 of *Mbl2*-201 (ENSMUST00000025797.7) transcript is recommended as the knockout region. The region contains 443 bp of coding sequences. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Mbl2* 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 on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.
- 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.



Gene Information





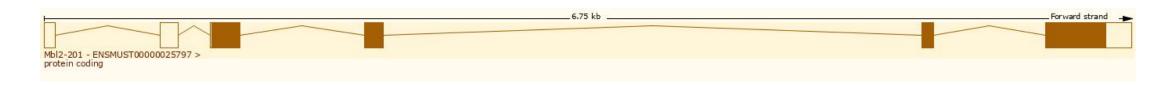


Transcript Information

The gene has 1 transcript, the transcript are shown below:

Transcript ID 🔻		Name 🝦	bp 👙	Protein 🝦	Biotype 🝦	CCDS	UniProt Match	Flags			
ENSMUST00000025797.7	7 1	Mbl2-201	1089	<u>244aa</u>	Protein coding	<u>CCDS29742</u> &	P41317@ Q3UEK1@	Ensembl Canonical	GENCODE basic	APPRIS P1	TSL:1

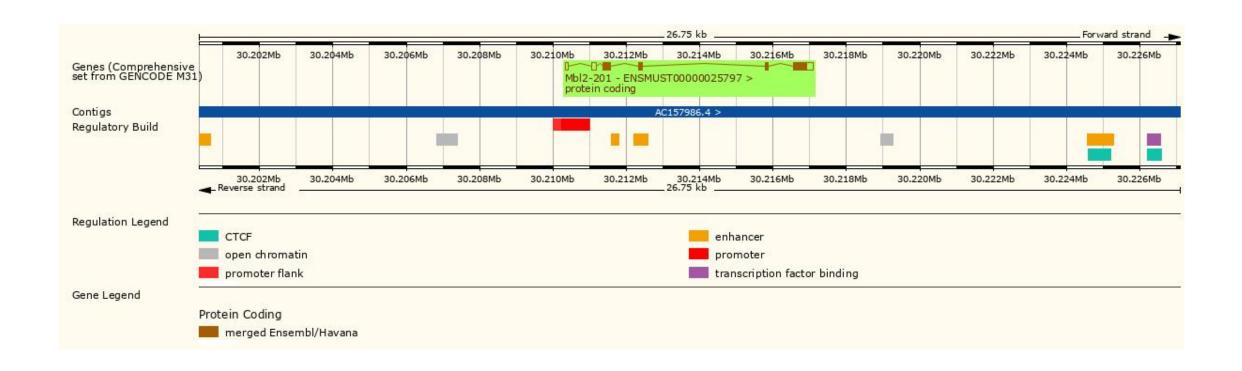
The strategy is based on the design of *Mbl2*-201 transcript, the transcription is shown below:



Source: https://www.ensembl.org



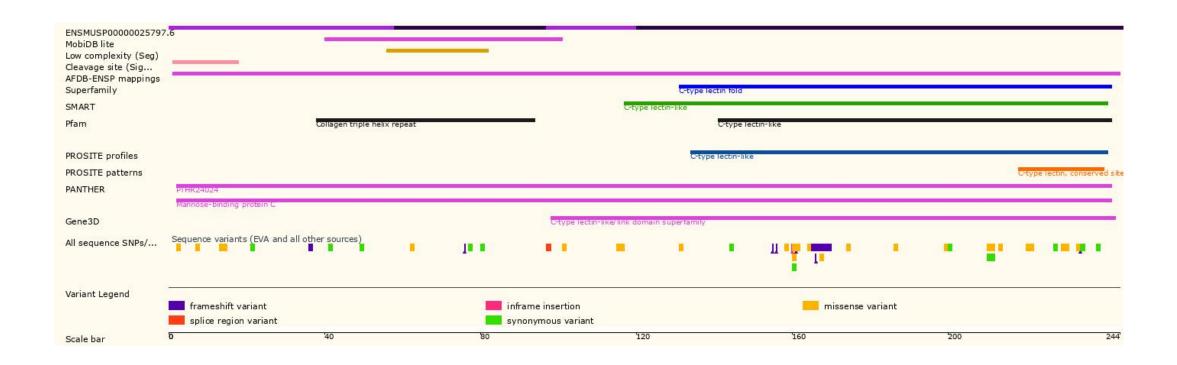
Genomic Information





Source: : https://www.ensembl.org

Protein Information





Source: : https://www.ensembl.org

Important Information

- A part of amino acid sequence will still remain at the N-terminal of the Mbl2 gene.
- *Mbl2* is located on Chr 19. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is 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 the existing technology level.

