

Lactb Cas9-CKO Strategy

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Overview

Target Gene Name

• Lactb

Project Type

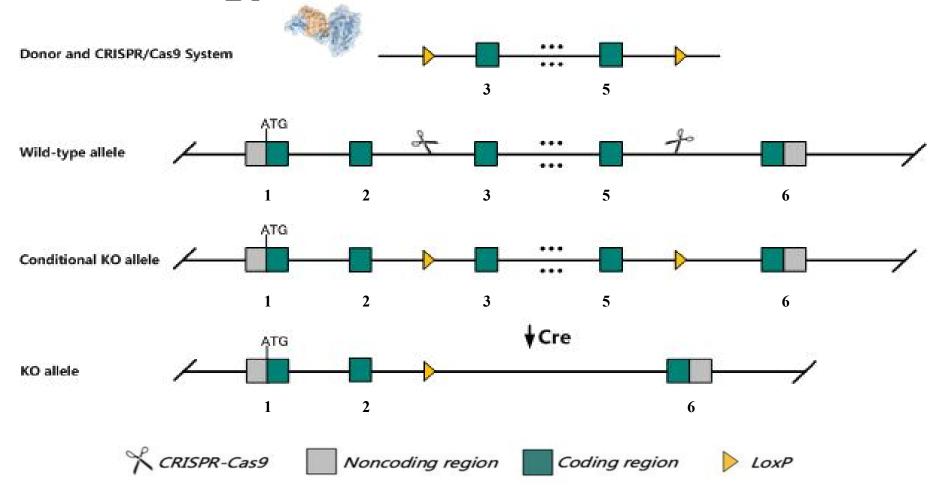
• Cas9-CKO

Genetic Background

• C57BL/6JGpt



Strain Strategy

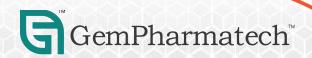


Schematic representation of CRISPR-Cas9 engineering used to edit the Lactb gene.



Technical Information

- The *Lactb* gene has 2 transcripts. According to the structure of *Lactb* gene, exon3-exon5 of *Lactb*-201 (ENSMUST00000034929.7) transcript is recommended as the knockout region. The region contains 712bp coding sequence. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Lactb* 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

Lactb lactamase, beta [Mus musculus (house mouse)]

≛ Download Datasets

Gene ID: 80907, updated on 5-Mar-2024



☆ ?

Official Symbol Lactb provided by MGI

Official Full Name lactamase, beta provided by MGI

Primary source MGI:MGI:1933395

See related Ensembl: ENSMUSG00000032370 AllianceGenome: MGI:1933395

Gene type protein coding RefSeg 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 Lact1; LACT-1; Mrpl56

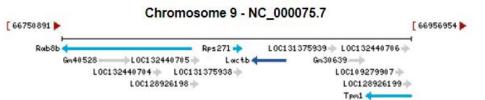
Summary Predicted to enable identical protein binding activity and peptidase activity. Predicted to be involved in proteolysis and regulation of lipid metabolic process. Predicted

to act upstream of or within lipid metabolic process. Located in mitochondrion. Is expressed in several structures, including early conceptus; gonad; heart; liver; and

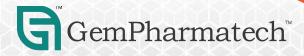
metanephros. Orthologous to human LACTB (lactamase beta), [provided by Alliance of Genome Resources, Apr 2022]

Expression Ubiquitous expression in duodenum adult (RPKM 13.5), adrenal adult (RPKM 10.4) and 28 other tissues See more

Orthologs human all



Source: https://www.ncbi.nlm.nih.gov/

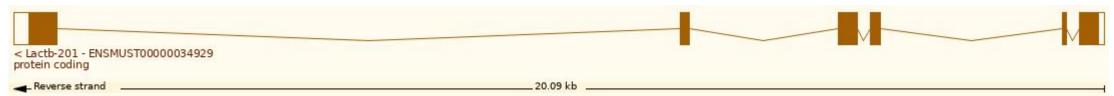


Transcript Information

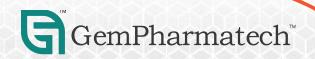
The gene has 2 transcripts, all transcripts are shown below:

Transcript ID	Name 🍦	bp 🌲	Protein	Biotype	CCDS 🍦	UniProt Match	Flags			
ENSMUST00000034929.7	Lactb-201	2031	<u>551aa</u>	Protein coding	CCDS40674 ₺	B2RWI2 छ Q9EP89 छ	Ensembl Canonical	GENCODE basic	APPRIS P1	TSL:1
ENSMUST00000215172.2	Lactb-202	1806	356aa	Protein coding		A0A1L1SVF9┏	GENCODE basic TSL:1			

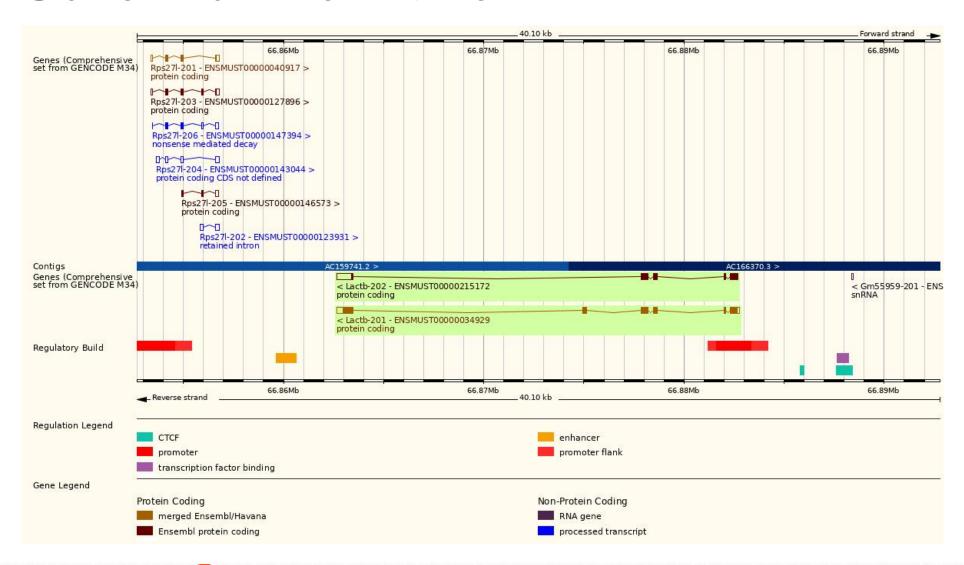
The strategy is based on the design of *Lactb*-201 transcript, the transcription is shown below:



Source: https://www.ensembl.org



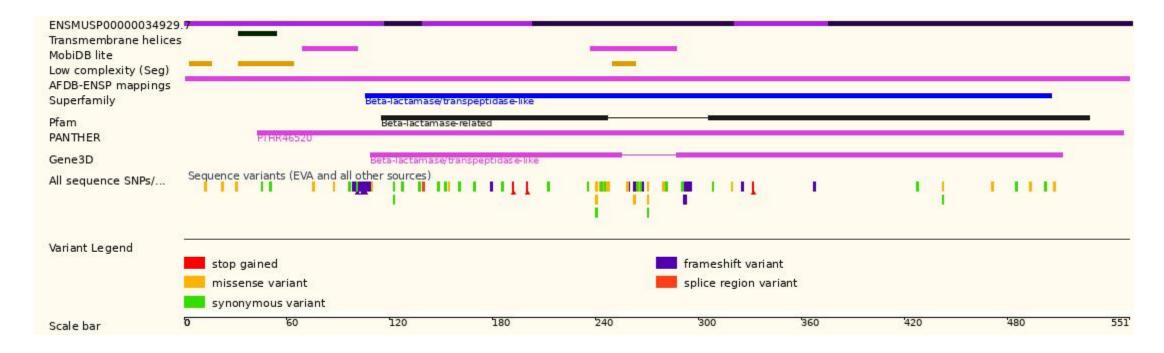
Genomic Information

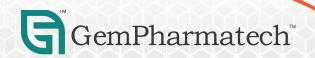




Source: : https://www.ensembl.org

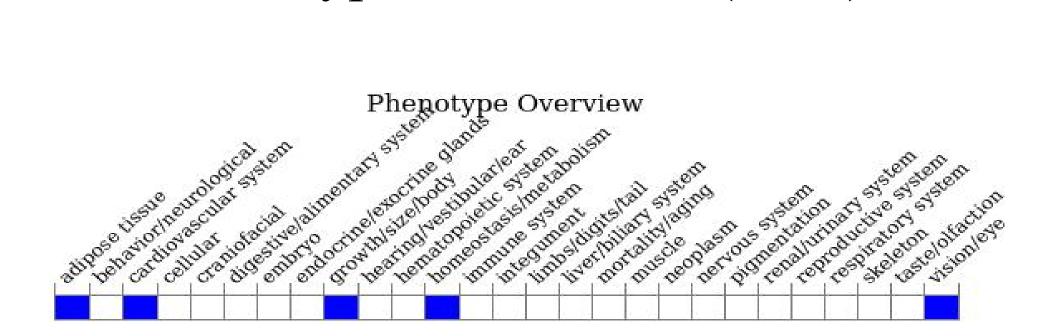
Protein Information





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

Mouse Phenotype Information (MGI)





Source: https://www.informatics.jax.org

Important Information

- This strategy does not cause frameshift mutation for *Lactb*-202, and it may leave some protein structures, the retained function is unknown.
- *Lactb* is located on Chr9. 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.

