

Cpb2 Cas9-KO Strategy

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Design Date: 2024-1-23

Overview

Target Gene Name

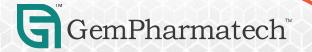
• Cpb2

Project Type

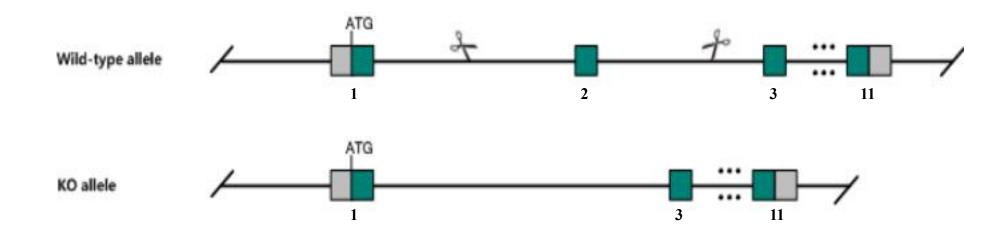
• Cas9-KO

Genetic Background

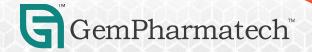
• C57BL/6JGpt



Strain Strategy







Technical Information

- The *Cpb2* gene has 2 transcripts. According to the structure of *Cpb2* gene, exon2 of *Cpb2*-201 (ENSMUST00000022576.10) transcript is recommended as the knockout region. The region contains 76bp coding sequence. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Cpb2* gene. The brief process is as follows: gRNAs were transcribed in vitro. Cas9 and gRNAs 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 ontarget 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.



Gene Information

Cpb2 carboxypeptidase B2 [Mus musculus (house mouse)]

Gene ID: 56373, updated on 22-Jan-2024



Official Symbol Cpb2 provided by MGI

Official Full Name carboxypeptidase B2 provided by MGI

Primary source MGI:MGI:1891837

See related Ensembl: ENSMUSG00000021999 Alliance Genome: MGI:1891837

Gene type protein coding RefSeq status REVIEWED

Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae;

Murinae; Mus; Mus

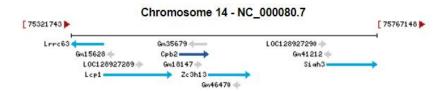
Also known as CPR; Cpu; TAFI; 1110032P04Rik; 4930405E17Rik

Summary This gene encodes carboxypeptidase B, a zinc-dependent metalloprotease that cleaves peptide bonds at the C-terminus of protein substrates. The encoded

preproprotein undergoes proteolytic activation to generate a mature, functional enzyme, and secreted into plasma. [provided by RefSeq, Jan 2016]

Expression Biased expression in liver E18 (RPKM 213.1), liver adult (RPKM 49.9) and 2 other tissues See more

Orthologs <u>human</u> all



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

≛ Download Datasets



Transcript Information

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

Transcript ID 🗼	Name ▲	bp 🌲	Protein 🍦	Biotype	CCDS .	UniProt Match	Flags			
ENSMUST00000022576.10	Cpb2-201	1477	<u>422aa</u>	Protein coding	CCDS27277 ₺	Q9JHH6&	Ensembl Canonical	GENCODE basic	APPRIS P1	TSL:1
ENSMUST00000227817.2	Cpb2-202	853	No protein	Retained intron		=		2		

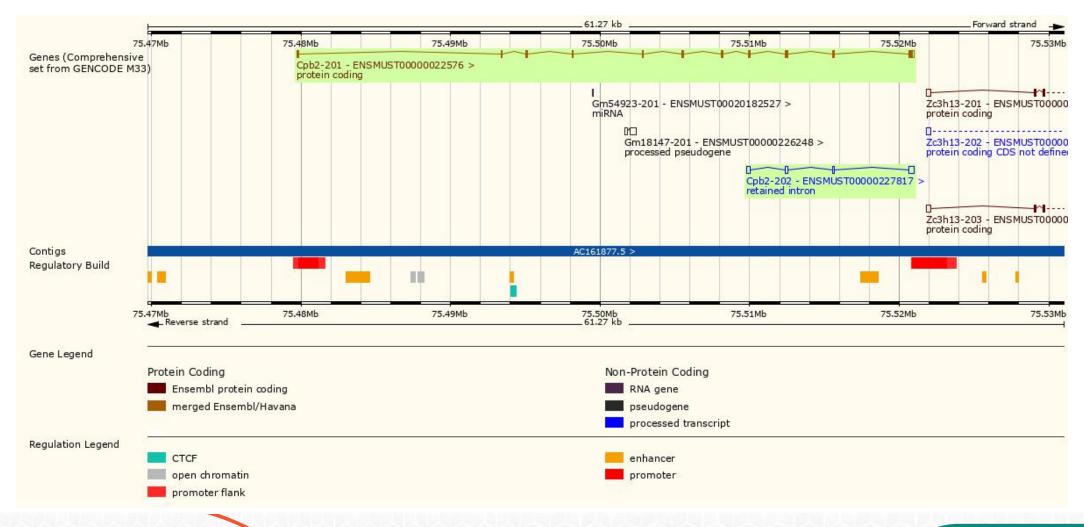
The strategy is based on the design of *Cpb2*-201 transcript, the transcription is shown below:

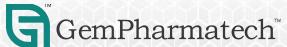


Source: https://www.ensembl.org



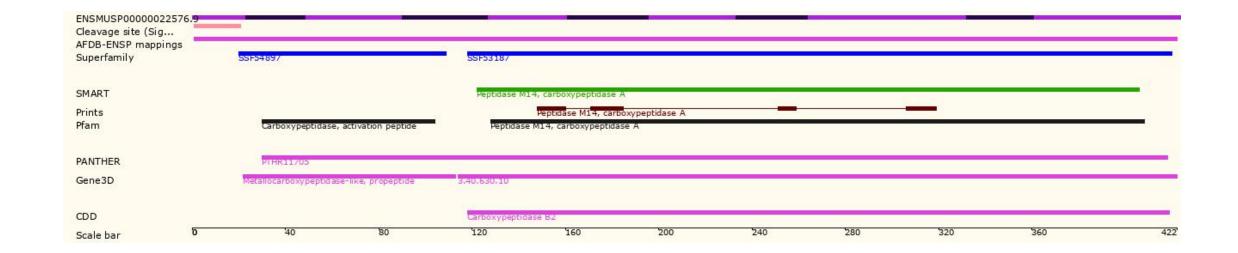
Genomic Information





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

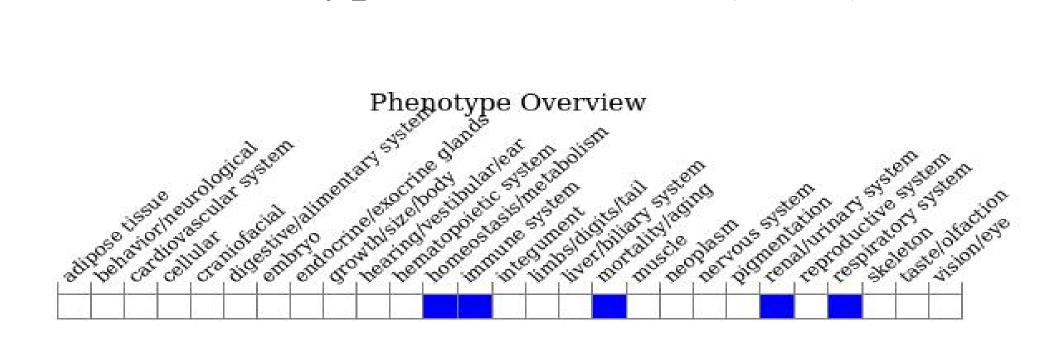
Protein Information





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

Mouse Phenotype Information (MGI)



• Homozygous null mice exhibit altered plasma clot lysis and may show reduced bleomycin-induced lung fibrosis, impaired healing of cutaneous wounds and colonic anastomoses, altered glomerular structure, or complement-mediated lethal inflammation after LPS sensitization.



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

Important Information

- Based the data of MGI, knock out mice deleting Exons 4 and 5 showed prenatal lethality, considerably reduced numbers of homozygotes are born from matings of heterozygotes.
- *Cpb2* is located on Chr14. 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 risks of the mutation on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

