

Car2 Cas9-KO Strategy

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Date:2019-10-27

Project Overview



Project Name Car2

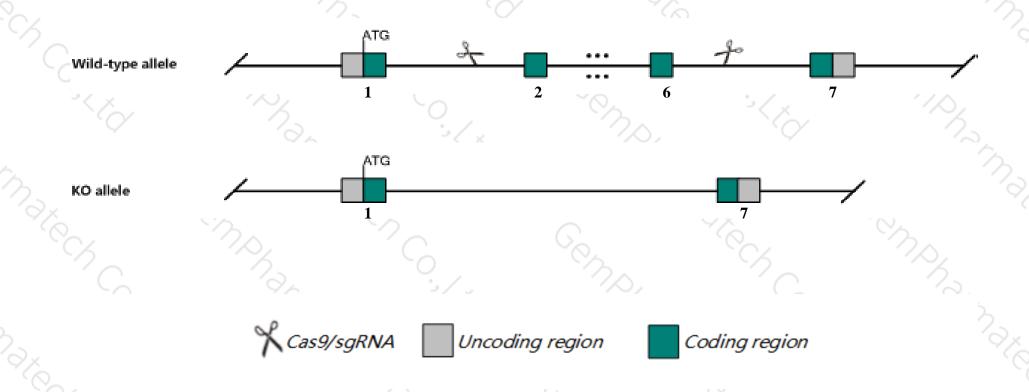
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Car2 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Car2* gene has 3 transcripts. According to the structure of *Car2* gene, exon2-exon6 of *Car2-201* (ENSMUST00000029078.8) transcript is recommended as the knockout region. The region contains 629bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Car2* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA 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.

Notice



- ➤ According to the existing MGI data, Homozygous mutant mice are growth retarded, display renal tubular acidosis, but mutants have not been recovered that display osteopetrosis as found in human CA-II deficiency.
- The *Car2* gene is located on the Chr3. 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 gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Car2 carbonic anhydrase 2 [Mus musculus (house mouse)]

Gene ID: 12349, updated on 24-Oct-2019

Summary

2 7

Official Symbol Car2 provided by MGI

Official Full Name carbonic anhydrase 2 provided by MGI

Primary source MGI:MGI:88269

See related Ensembl:ENSMUSG00000027562

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

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

Muridae; Murinae; Mus; Mus

Also known as Ca2; CAII; Car-2; Ltw-5; Lvtw-5; Al131712

Expression Biased expression in liver E14 (RPKM 556.3), liver E14.5 (RPKM 538.2) and 14 other tissues See more

Orthologs <u>human</u> all

Genomic context



Location: 3 A1; 3 3.23 cM

See Car2 in Genome Data Viewer

Exon count: 7

| Annotation release | Status | Assembly | Chr | Location | |
|--------------------|-------------------|------------------------------|-----|--------------------------------|--|
| 108 | current | GRCm38.p6 (GCF 000001635.26) | 3 | NC_000069.6 (1488626914900770) | |
| Build 37.2 | previous assembly | MGSCv37 (GCF 000001635.18) | 3 | NC_000069.5 (1488642614900770) | |

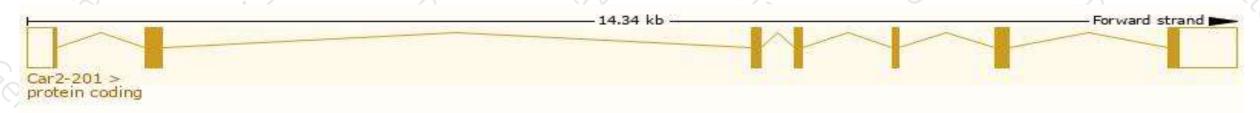
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

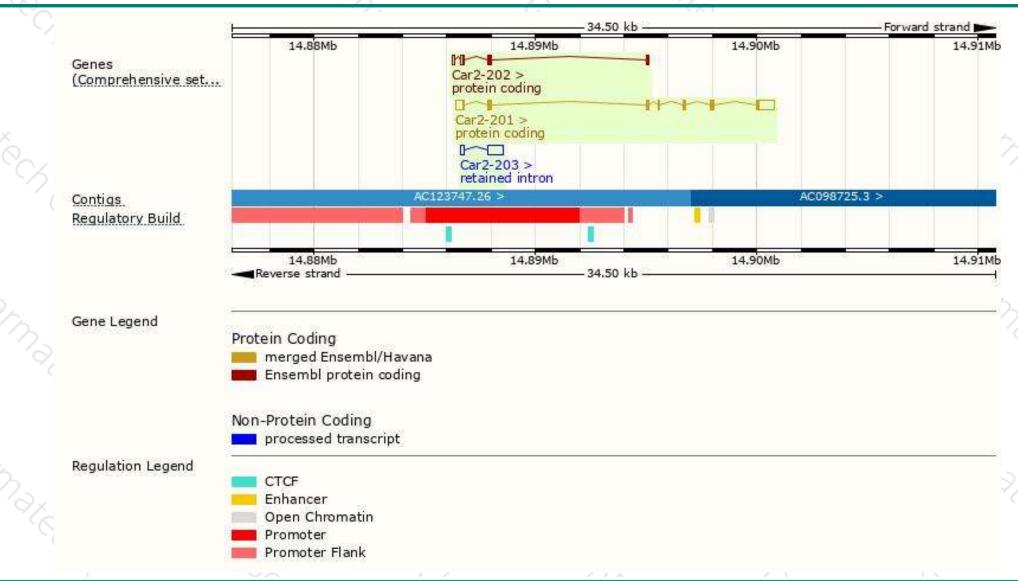
| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|----------|----------------------|------|--------------|-----------------|-----------|------------|-------------------------------|
| Car2-201 | ENSMUST00000029078.8 | 1788 | <u>260aa</u> | Protein coding | CCDS17251 | P00920 | TSL:1 GENCODE basic APPRIS P1 |
| Car2-202 | ENSMUST00000192609.5 | 587 | <u>115aa</u> | Protein coding | - | A0A0A6YX78 | CDS 3' incomplete TSL:3 |
| Car2-203 | ENSMUST00000195520.1 | 881 | No protein | Retained intron | - | - | TSL:2 |

The strategy is based on the design of Car2-201 transcript, The transcription is shown below



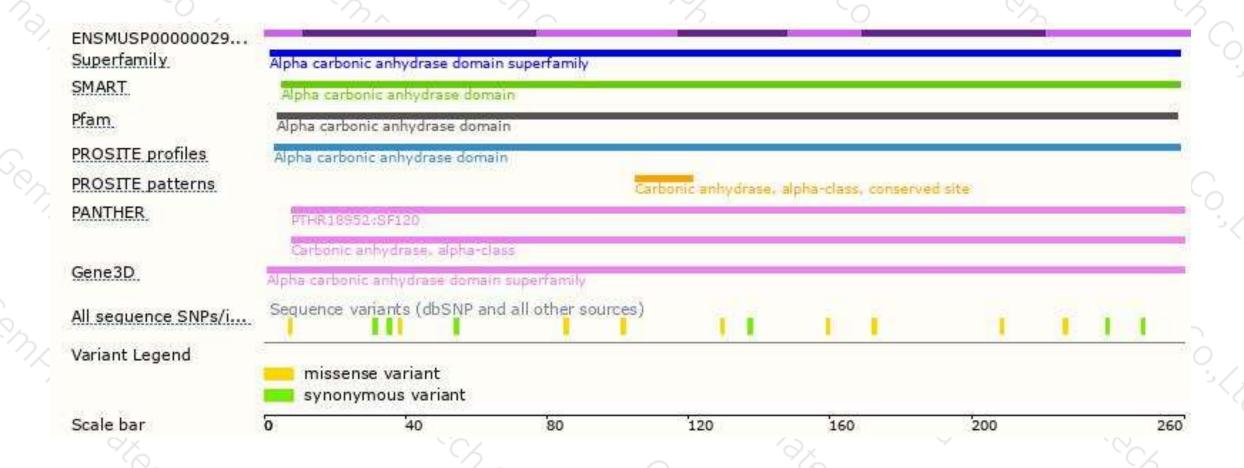
Genomic location distribution





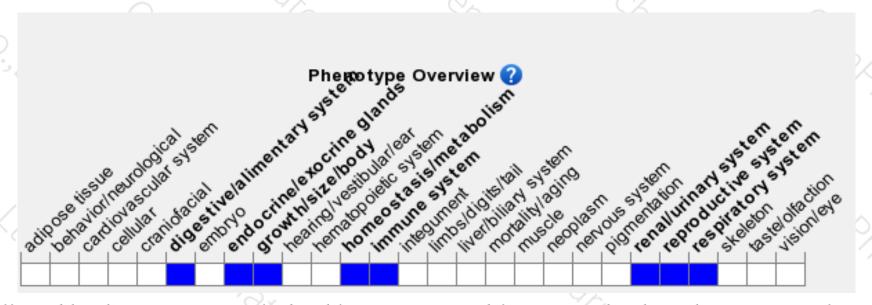
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(http://www.informatics.jax.org/).

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

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