

Gpr157 Cas9-KO Strategy

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Project Overview



Project Name

Gpr157

Project type

Cas9-KO

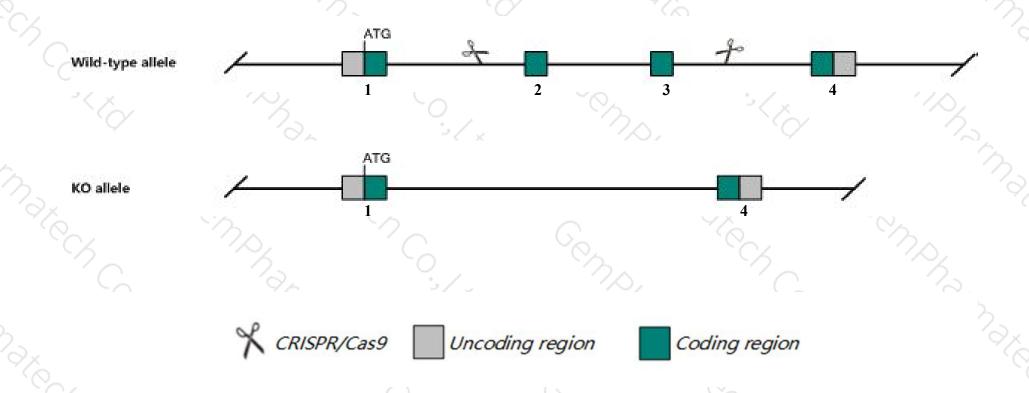
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gpr157* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Gpr157* gene has 1 transcript. According to the structure of *Gpr157* gene, exon2-exon3 of *Gpr157-201* (ENSMUST00000094451.3) transcript is recommended as the knockout region. The region contains 412bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Gpr157* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ The *Gpr157* gene is located on the Chr4. 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)



Gpr157 G protein-coupled receptor 157 [Mus musculus (house mouse)]

Gene ID: 269604, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Gpr157 provided by MGI

Official Full Name G protein-coupled receptor 157 provided by MGI

Primary source MGI:MGI:2442046

See related Ensembl:ENSMUSG00000047875

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 F730108M23Rik

Expression Broad expression in heart adult (RPKM 8.7), mammary gland adult (RPKM 3.0) and 22 other tissuesSee more

Orthologs human all

Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

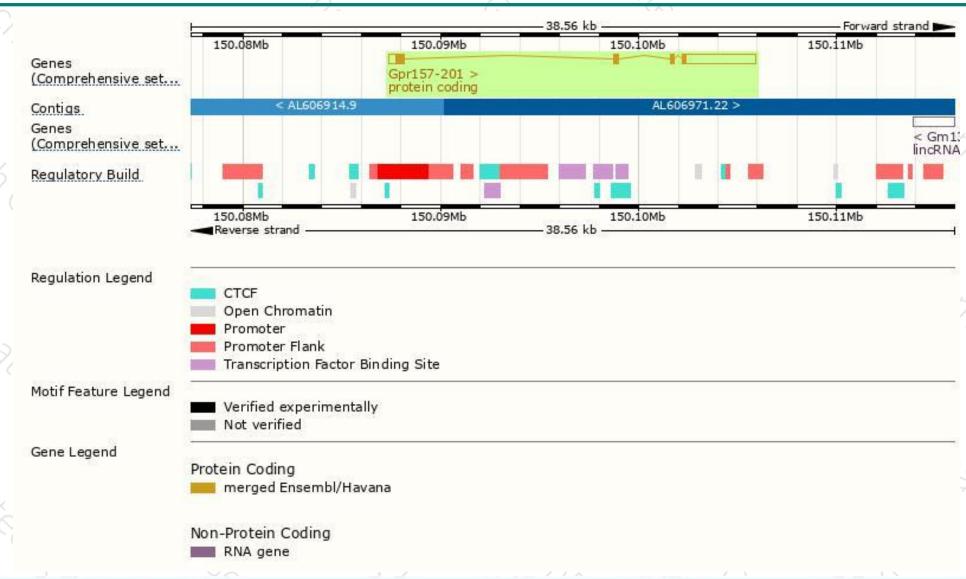
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Gpr157-201	ENSMUST00000094451.3	4900	330aa	Protein coding	CCDS38977	Q148S2 Q8C206	TSL:1 GENCODE basic APPRIS P1	L

The strategy is based on the design of Gpr157-201 transcript, The transcription is shown below



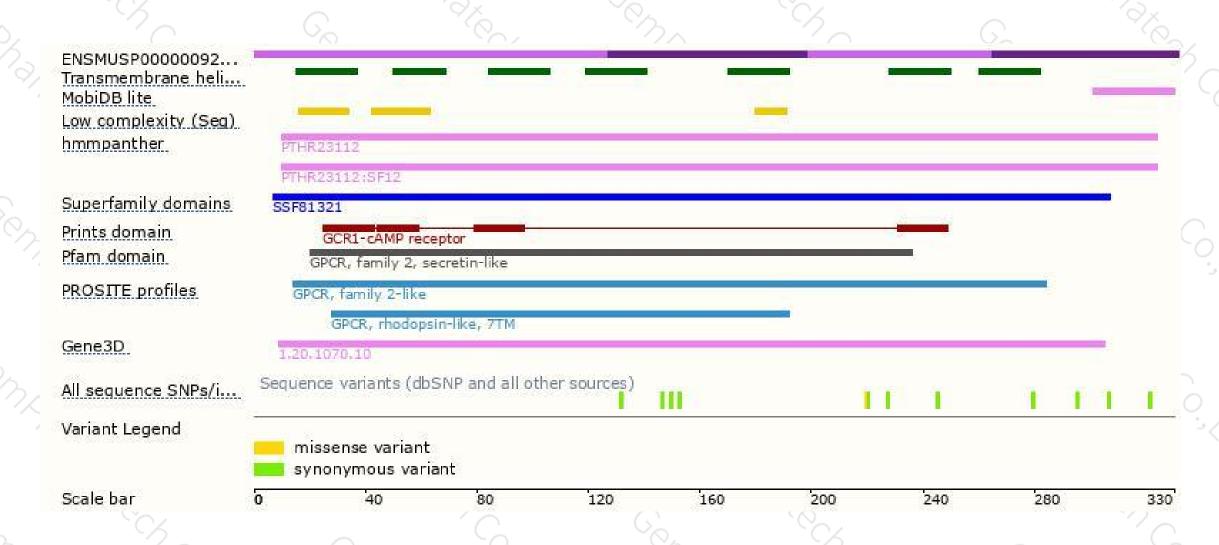
Genomic location distribution





Protein domain







If you have any questions, you are welcome to inquire. Tel: 400-9660890





