

Galr3 Cas9-KO Strategy

Designer:Xueting Zhang

Reviewer: Yanhua Shen

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Project Overview



Project Name

Galr3

Project type

Cas9-KO

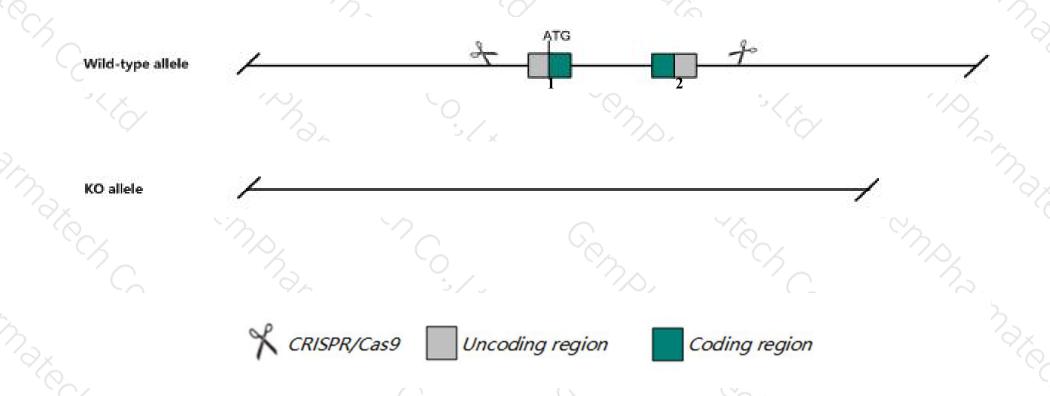
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Galr3 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Galr3* gene has 1 transcript. According to the structure of *Galr3* gene, exon1-exon2 of *Galr3-201* (ENSMUST0000058004.3) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Galr3* gene. The brief process is as follows: gRNA was transcribed in vitro.Cas9 and gRNA 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, mice homozygous for a null allele exhibit increased anxiety-like behavior and abnormal social investigation.
- The knockout region is near to the C-terminal of *Gm3924* gene, this strategy may influence the regulatory function of the C-terminal of *Gm3924* gene.
- > Gcat gene will be destroyed together in this strategy.
- The *Galr3* gene is located on the Chr15. 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)



Galr3 galanin receptor 3 [Mus musculus (house mouse)]

Gene ID: 14429, updated on 12-Aug-2019

Summary

☆ ?

Official Symbol Galr3 provided by MGI

Official Full Name galanin receptor 3 provided by MGI

Primary source MGI:MGI:1329003

See related Ensembl: ENSMUSG00000114755

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 Galnr3

Expression Biased expression in duodenum adult (RPKM 6.2), small intestine adult (RPKM 2.3) and 11 other tissues See more

Orthologs human all

Genomic context



Location: 15 E1; 15 37.7 cM

See Galr3 in Genome Data Viewer

Exon count: 2

| Annotation release | Status | Assembly | Chr | Location | |
|--------------------|-------------------|------------------------------|-----|--------------------------------|--|
| 108 | current | GRCm38.p6 (GCF_000001635.26) | 15 | NC_000081.6 (7904188579043558) | |
| Build 37.2 | previous assembly | MGSCv37 (GCF_000001635.18) | 15 | NC_000081.5 (7887231578873988) | |

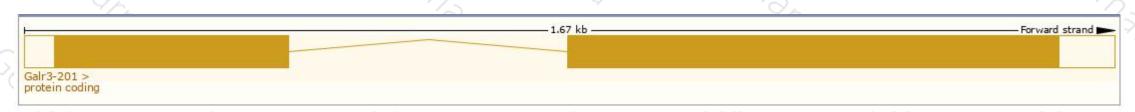
Transcript information (Ensembl)



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

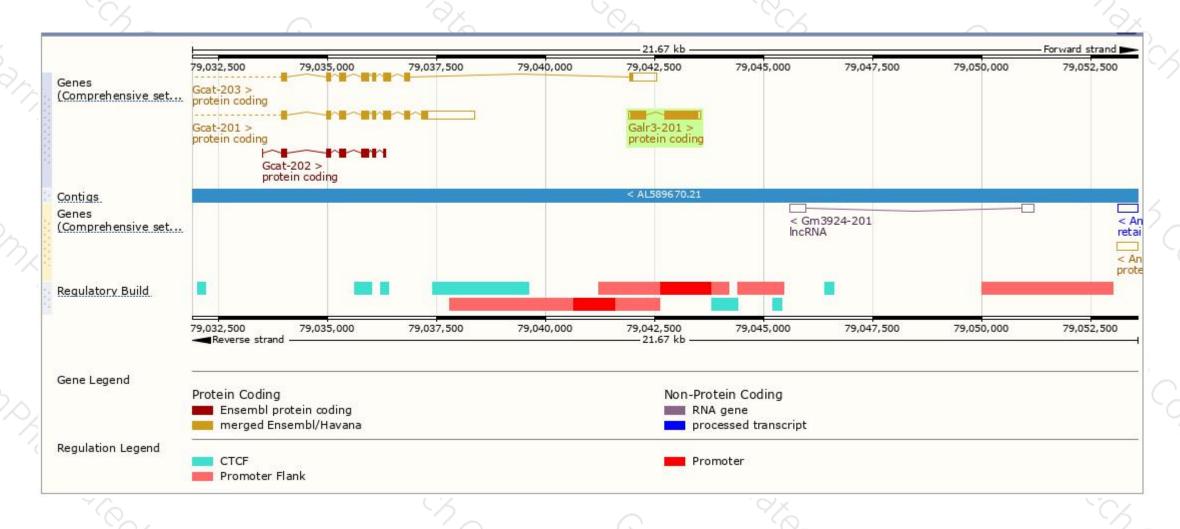
| Name 🍦 | Transcript ID 🍦 | bp 👙 | Protein 🌲 | Biotype | CCDS 🍦 | UniProt 🌲 | Flags | | |
|-----------|----------------------|------|--------------|----------------|-------------|-----------------|-------|---------------|-----------|
| Galr3-201 | ENSMUST00000058004.3 | 1245 | <u>370aa</u> | Protein coding | CCDS27631 ₽ | <u>088853</u> ₽ | TSL:1 | GENCODE basic | APPRIS P1 |

The strategy is based on the design of Galr3-201 transcript, The transcription is shown below



Genomic location distribution





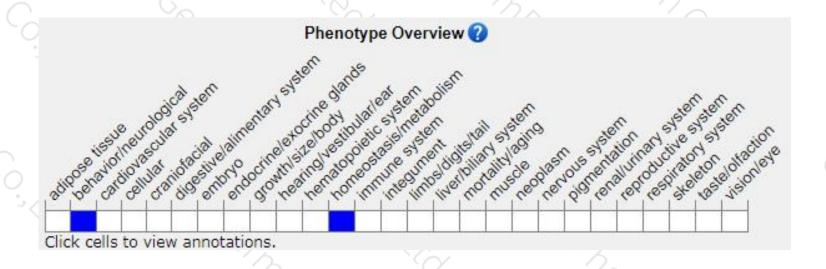
Protein domain





Mouse phenotype description(MGI)





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

According to the existing MGI data, Mice homozygous for a null allele exhibit increased anxiety-like behavior and abnormal social investigation.



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





