Gpr4 Cas9-KO Strategy

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Design Date:

2019-8-1

Project Overview



Project Name

Gpr4

Project type

Cas9-KO

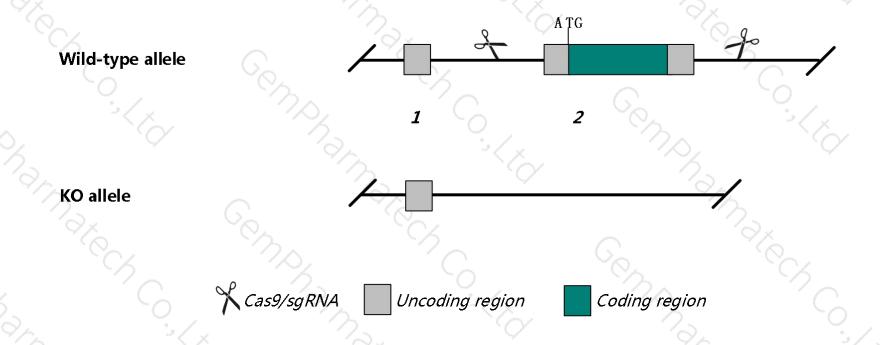
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gpr4* gene. The schematic diagram is as follows:



Technical routes



- The *Gpr4* gene has 1 transcripts. According to the structure of *Gpr4* gene, exon2 of *Gpr4*-201 (ENSMUST00000060225.5)transcript is recommended as the knockout region. The region contains 1098bp 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 *Gpr4* 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



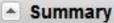
- According to the existing MGI data, mice homozygous for a null allele display partial neonatal and postnatal lethality, hemorrhages, impaired association of vascular smooth muscle cells with capillaries and small arteries and veins, and impaired contact between mesangial cells and renal glomerular capillaries.
- The xxx gene is located on the Chr7. 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)



Gpr4 G protein-coupled receptor 4 [Mus musculus (house mouse)]

Gene ID: 319197, updated on 8-Dec-2018





Official Symbol Gpr4 provided by MGI

Official Full Name G protein-coupled receptor 4 provided by MGI

Primary source MGI:MGI:2441992

See related Ensembl: ENSMUSG00000044317

RefSeq status VALIDATED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria;

Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Expression Ubiquitous expression in lung adult (RPKM 11.4), subcutaneous fat pad adult (RPKM 11.1) and 25

other tissues See more

Orthologs human all

Transcript information (Ensembl)



The gene has 1 transcript, and all transcripts are shown below:

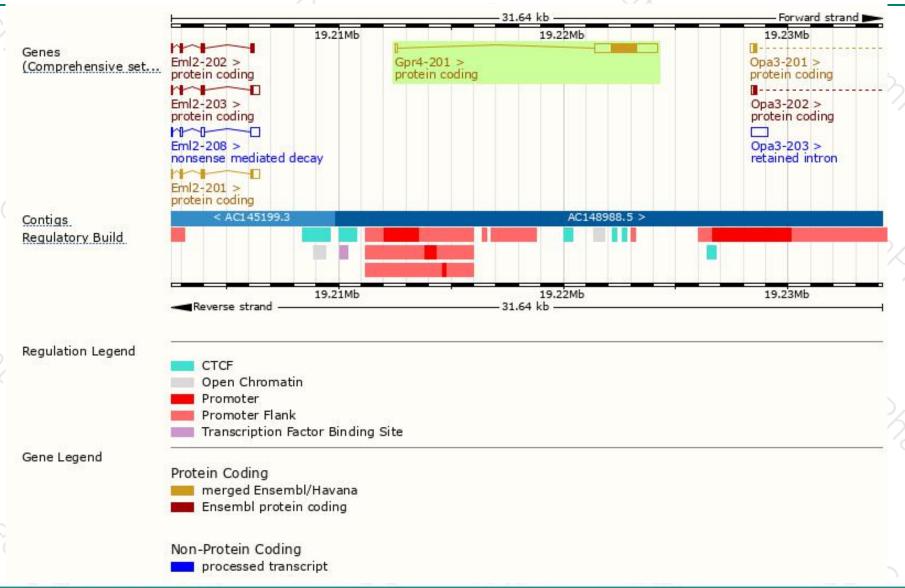
Show/hide columns (1 hidden)									
Name 🌲	Transcript ID	bp 🍦	Protein 4	Biotype	CCDS 🍦	UniProt #	RefSeq 🍦	Flags	
Gpr4-201	ENSMUST00000060225.5	2882	<u>365aa</u>	Protein coding	CCDS20893&	Q8BUD0@	NM 175668 & NP 783599 &	TSL:1 GENCODE basic	APPRIS P1

The strategy is based on the design of *Gpr4*-201 transcript, The transcription is shown below



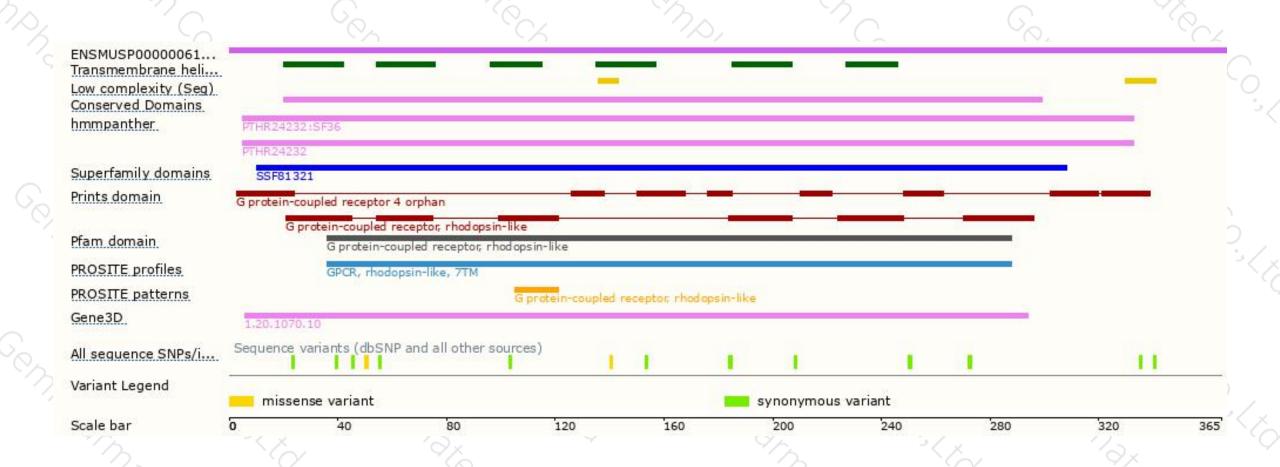
Genomic location distribution





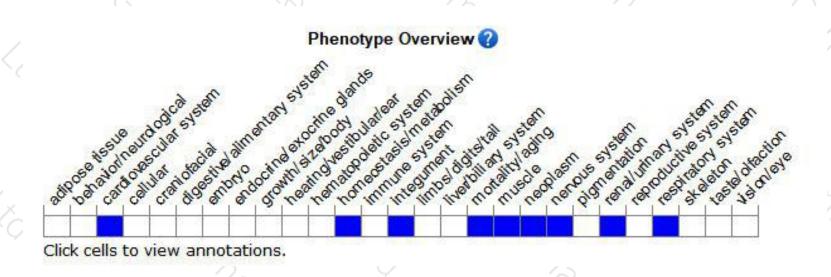
Protein domain





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 display partial neonatal and postnatal lethality, hemorrhages, impaired association of vascular smooth muscle cells with capillaries and small arteries and veins, and impaired contact between mesangial cells and renal glomerular capillaries.

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





