

Gpr83 Cas9-CKO Strategy

Designer:

Reviewer

Design Date:

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Project Overview



Project Name

Gpr83

Project type

Cas9-CKO

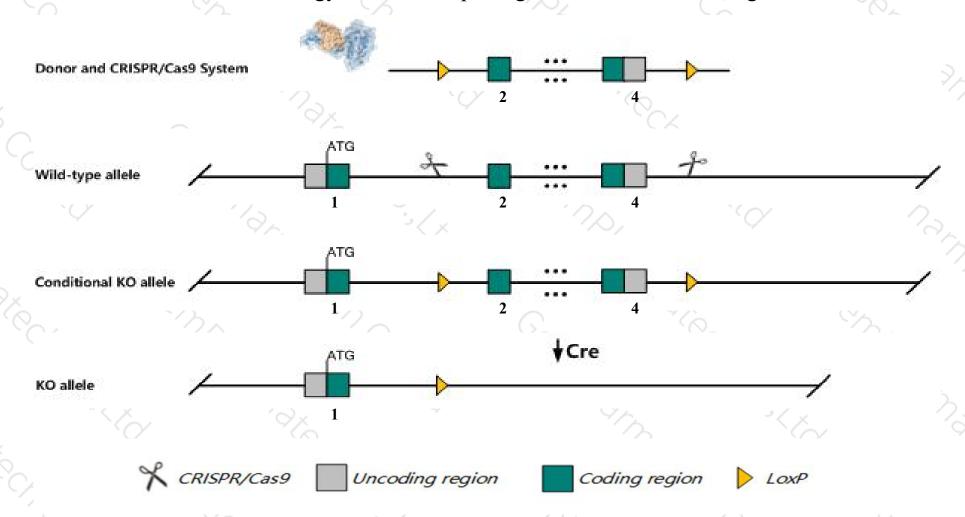
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gpr83* gene. The schematic diagram is as follows:



Technical routes



- The *Gpr83* gene has 3 transcripts. According to the structure of *Gpr83* gene, exon2-exon4 of *Gpr83-201* (ENSMUST00000034408.11) transcript is recommended as the knockout region. The region contains most 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 *Gpr83* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- ➤ According to the existing MGI data, mice homozygous for a null allele exhibit normal regulatory T cell development. Mice homozygous for a different targeted allele exhibit attenuated stress-evoked anxiety, increased sucrose preference and delayed spatial learning.
- > The *Gpr83* gene is located on the Chr9. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Gpr83 G protein-coupled receptor 83 [Mus musculus (house mouse)]

Gene ID: 14608, updated on 12-Aug-2019

Summary

☆ ?

Official Symbol Gpr83 provided by MGI

Official Full Name G protein-coupled receptor 83 provided by MGI

Primary source MGI:MGI:95712

See related Ensembl: ENSMUSG00000031932

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 Gir; RP39; RP82; Gpr72; RP105; AW045253

Expression Biased expression in cortex adult (RPKM 4.8), frontal lobe adult (RPKM 3.6) and 11 other tissues See more

Orthologs <u>human</u> all

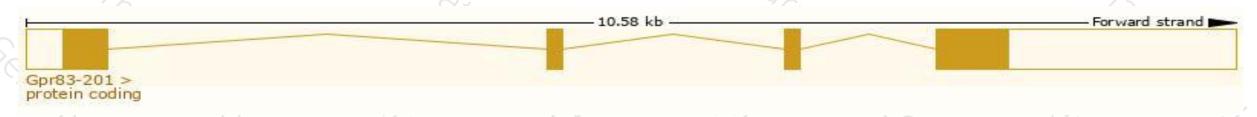
Transcript information (Ensembl)



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

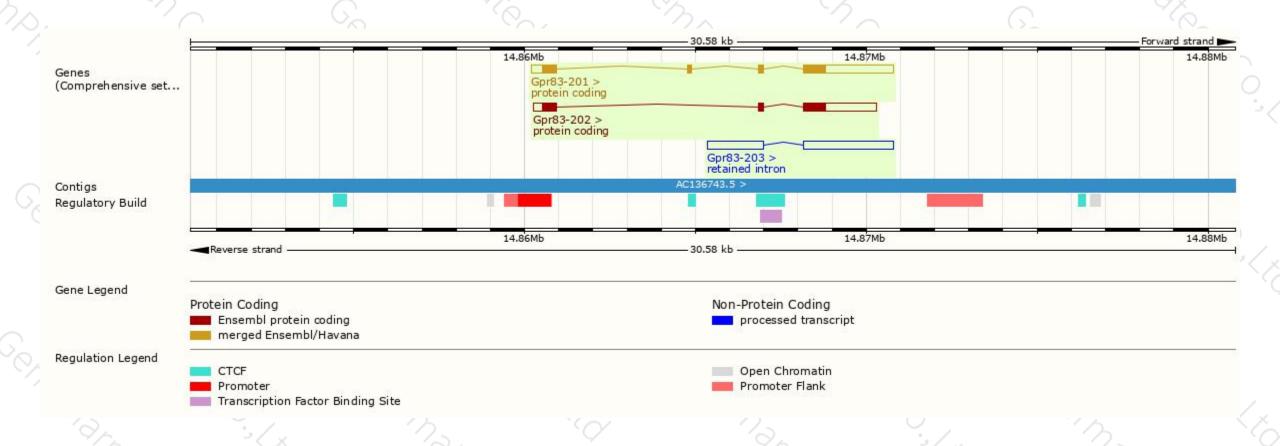
Name	Transcript ID 🔺	bp 🛊	Protein	Biotype	CCDS	UniProt	Flags
Gpr83-201	ENSMUST00000034408.11	3586	423aa	Protein coding	CCDS22829₺	P30731&Q542Q9&	TSL:1 GENCODE basic APPRIS P1
Gpr83-202	ENSMUST00000115624.2	2937	<u>381aa</u>	Protein coding	CCDS80959₽	P30731& Q544C4&	TSL:1 GENCODE basic
Gpr83-203	ENSMUST00000133278.1	4258	No protein	Retained intron	5	151	TSL:1

The strategy is based on the design of *Gpr83-201* transcript, The transcription is shown below



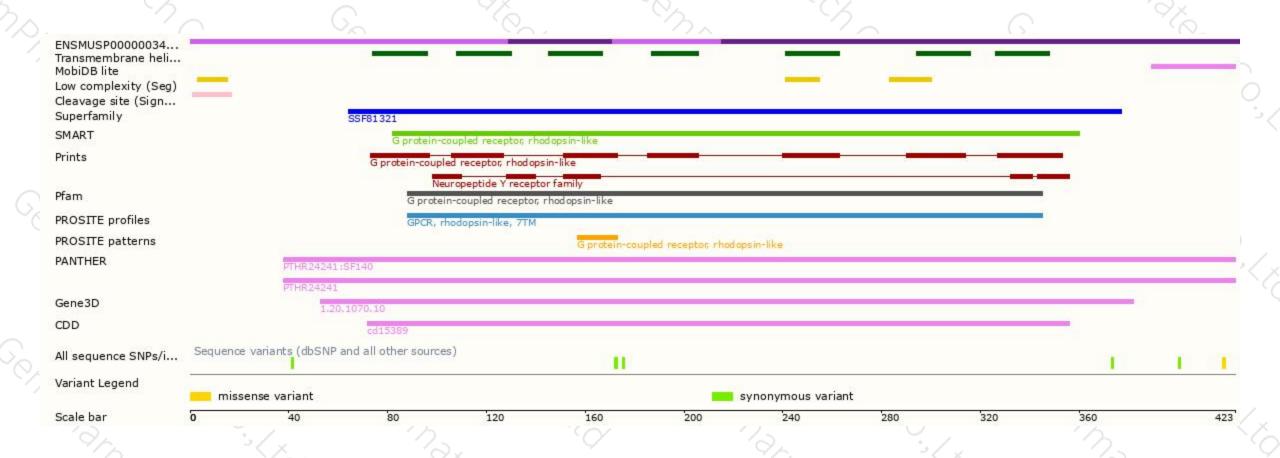
Genomic location distribution





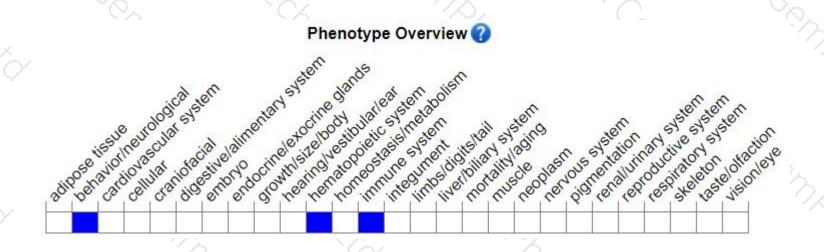
Protein domain





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 normal regulatory T cell development. Mice homozygous for a different targeted allele exhibit attenuated stress-evoked anxiety, increased sucrose preference and delayed spatial learning.



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





