

Gpr143 Cas9-CKO Strategy

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Project Overview



Project Name

Gpr143

Project type

Cas9-CKO

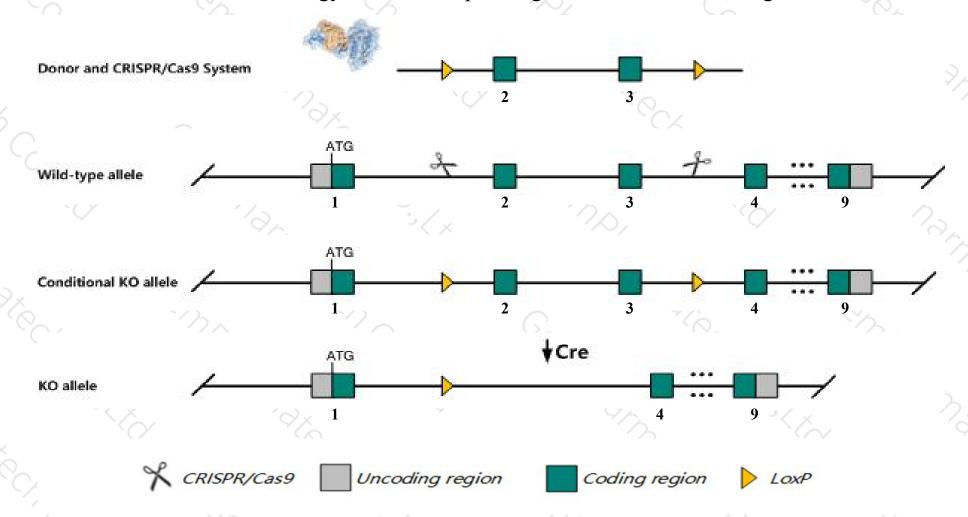
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gpr143* gene. The schematic diagram is as follows:



Technical routes



- The *Gpr143* gene has 3 transcripts. According to the structure of *Gpr143* gene, exon2-exon3 of *Gpr143-201* (ENSMUST00000026383.3) transcript is recommended as the knockout region. The region contains 205bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Gpr143* gene. The brief process is as follows:gRNA was transcribed in vitro, donor was constructed.Cas9, gRNA 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, Hemizygous males exhibit hypopigmentation of the ocular fundus, misrouting of the optic fibers at the chiasm, and the presence of giant melanosomes in the pigment epithelium of the eye.
- The *Gpr143* gene is located on the ChrX. 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)



Gpr143 G protein-coupled receptor 143 [Mus musculus (house mouse)]

Gene ID: 18241, updated on 18-Apr-2019

Summary

Official Symbol Gpr143 provided by MGI

Official Full Name G protein-coupled receptor 143 provided by MGI

Primary source MGI:MGI:107193

See related Ensembl: ENSMUSG00000025333

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 Oa1

Expression Low expression observed in reference dataset See more

Orthologs human all

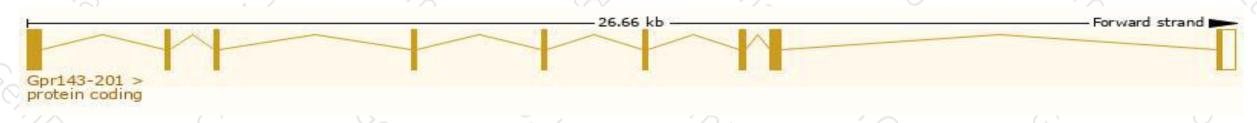
Transcript information (Ensembl)



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

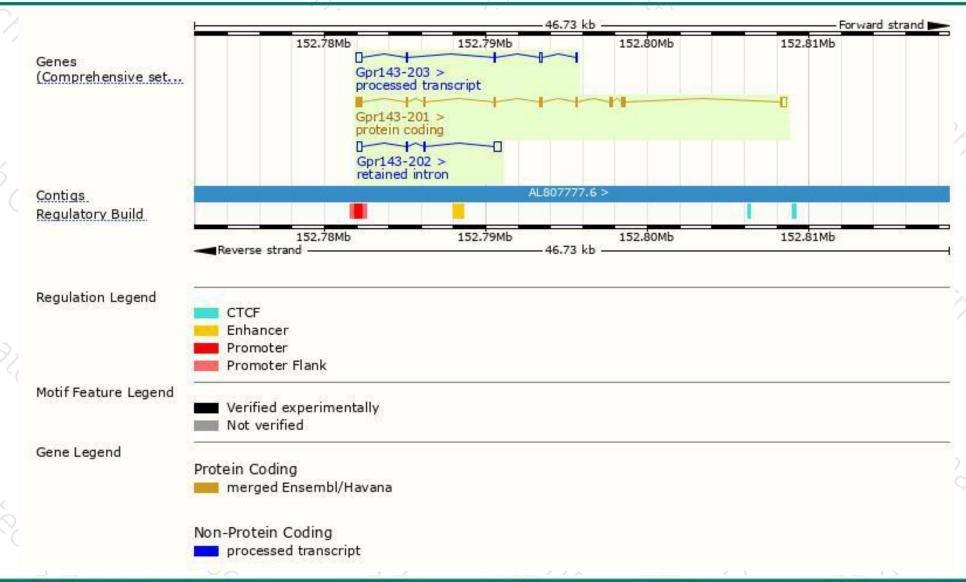
| Show/hide columns (1 hidden) | | | | | | | Filter |
|------------------------------|----------------------|------|------------|-----------------|------------|-----------------|-------------------------------|
| Name 🍦 | Transcript ID 🔻 | bp 🍦 | Protein | Biotype | CCDS 👙 | UniProt | Flags |
| Gpr143-203 | ENSMUST00000151819.7 | 745 | No protein | IncRNA | - | 2 | TSL:5 |
| Gpr143-202 | ENSMUST00000139310.1 | 917 | No protein | Retained intron | 125 | 10 | TSL:1 |
| Gpr143-201 | ENSMUST00000026383.3 | 1586 | 405aa | Protein coding | CCDS30476世 | P70259& Q549B6& | TSL:1 GENCODE basic APPRIS P1 |

The strategy is based on the design of *Gpr143-201* transcript, The transcription is shown below



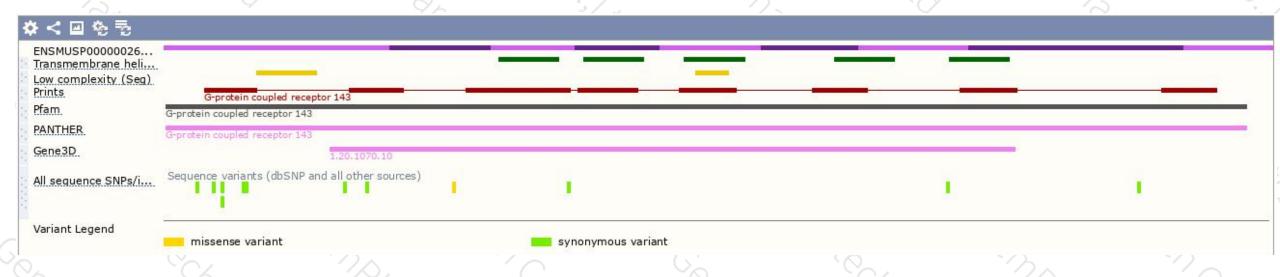
Genomic location distribution





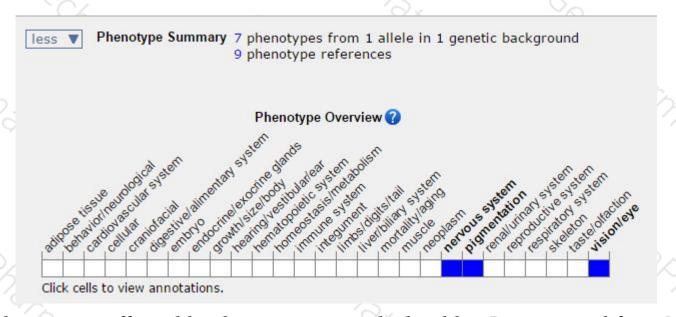
Protein domain





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, Hemizygous males exhibit hypopigmentation of the ocular fundus, misrouting of the optic fibers at the chiasm, and the presence of giant melanosomes in the pigment epithelium of the eye.



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





