



Gpr156 Cas9-CKO Strategy

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Project Overview

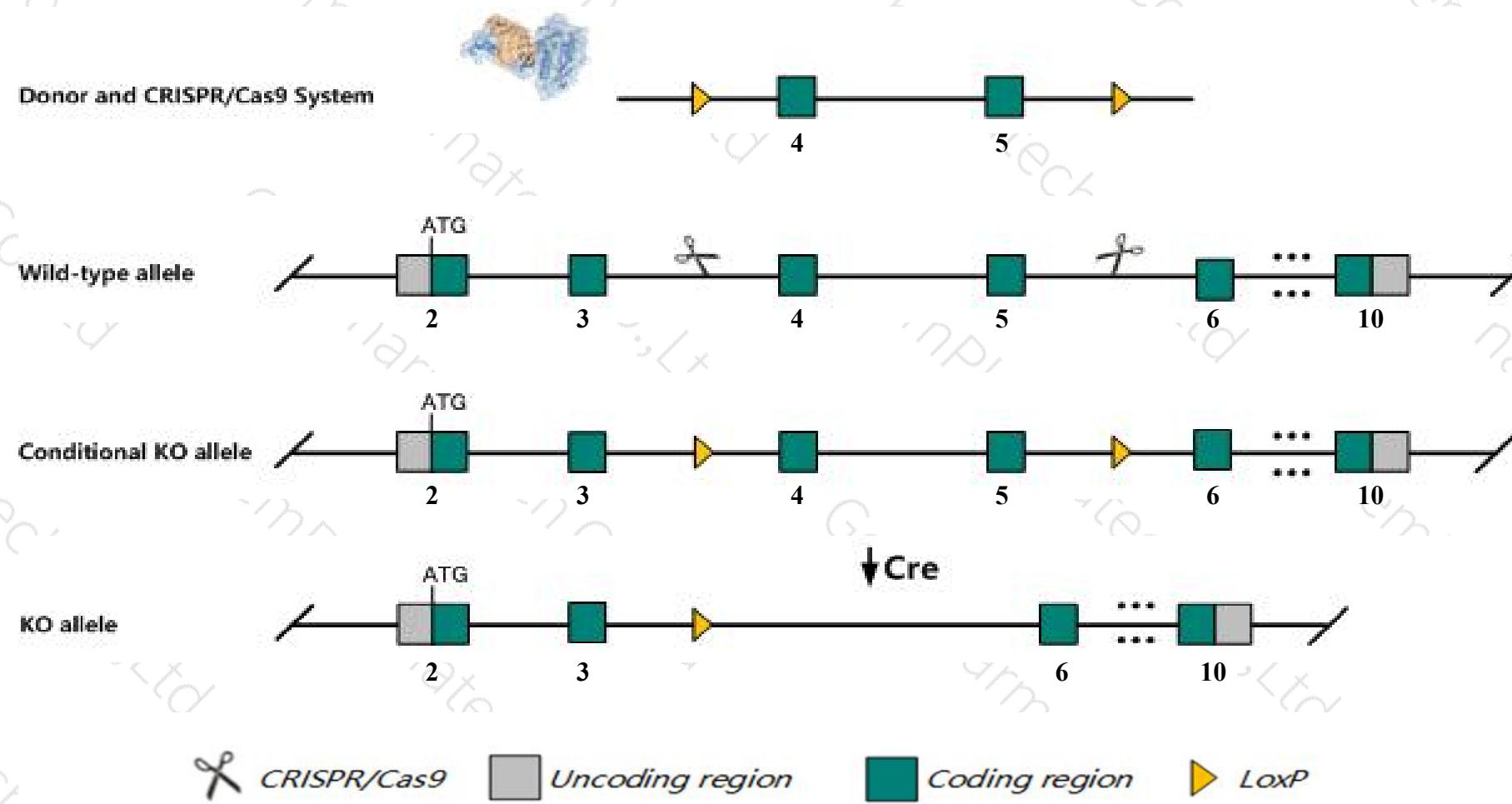
Project Name**Gpr156**

Project type**Cas9-CKO**

Strain background**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gpr156* gene. The schematic diagram is as follows:



Technical routes

- The *Gpr156* gene has 1 transcript. According to the structure of *Gpr156* gene, exon4-exon5 of *Gpr156-201* (ENSMUST00000061274.8) transcript is recommended as the knockout region. The region contains 241bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gpr156* 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.



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Notice

- The *Gpr156* gene is located on the Chr16. 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.



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Gene information (NCBI)

Gpr156 G protein-coupled receptor 156 [Mus musculus (house mouse)]

Gene ID: 239845, updated on 31-Jan-2019

Summary



Official Symbol Gpr156 provided by [MGI](#)

Official Full Name G protein-coupled receptor 156 provided by [MGI](#)

Primary source [MGI:MGI:2653880](#)

See related [Ensembl:ENSMUSG00000046961](#)

Gene type protein coding

RefSeq status PROVISIONAL

Organism [Mus musculus](#)

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as Gabab1

Expression Biased expression in CNS E14 (RPKM 1.9), whole brain E14.5 (RPKM 1.9) and 13 other tissues [See 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
Gpr156-201	ENSMUST00000061274.8	4584	798aa	Protein coding	CCDS28162	A0A0R4J0P3	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Gpr156-201* transcript, The transcription is shown below



Genomic location distribution



Gene Legend

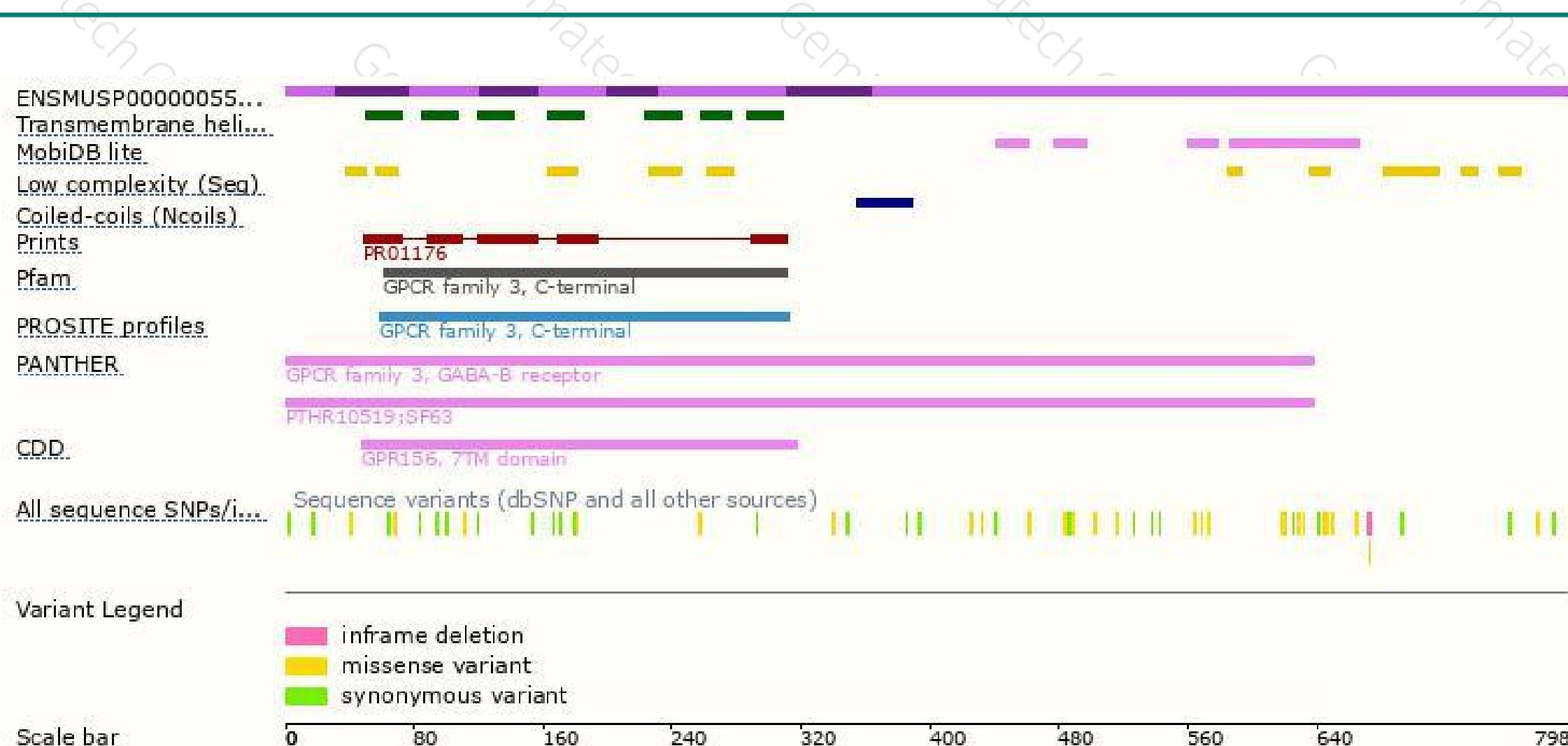
Protein Coding
merged Ensembl/Havana

Regulation Legend

Non-Protein Coding
RNA gene

CTCF
Enhancer
Open Chromatin
Promoter
Promoter Flank
Transcription Factor Binding Site

Protein domain





If you have any questions, you are welcome to inquire.

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