

# ***Gpr52* Cas9-CKO Strategy**

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# Project Overview

**Project Name**

***Gpr52***

**Project type**

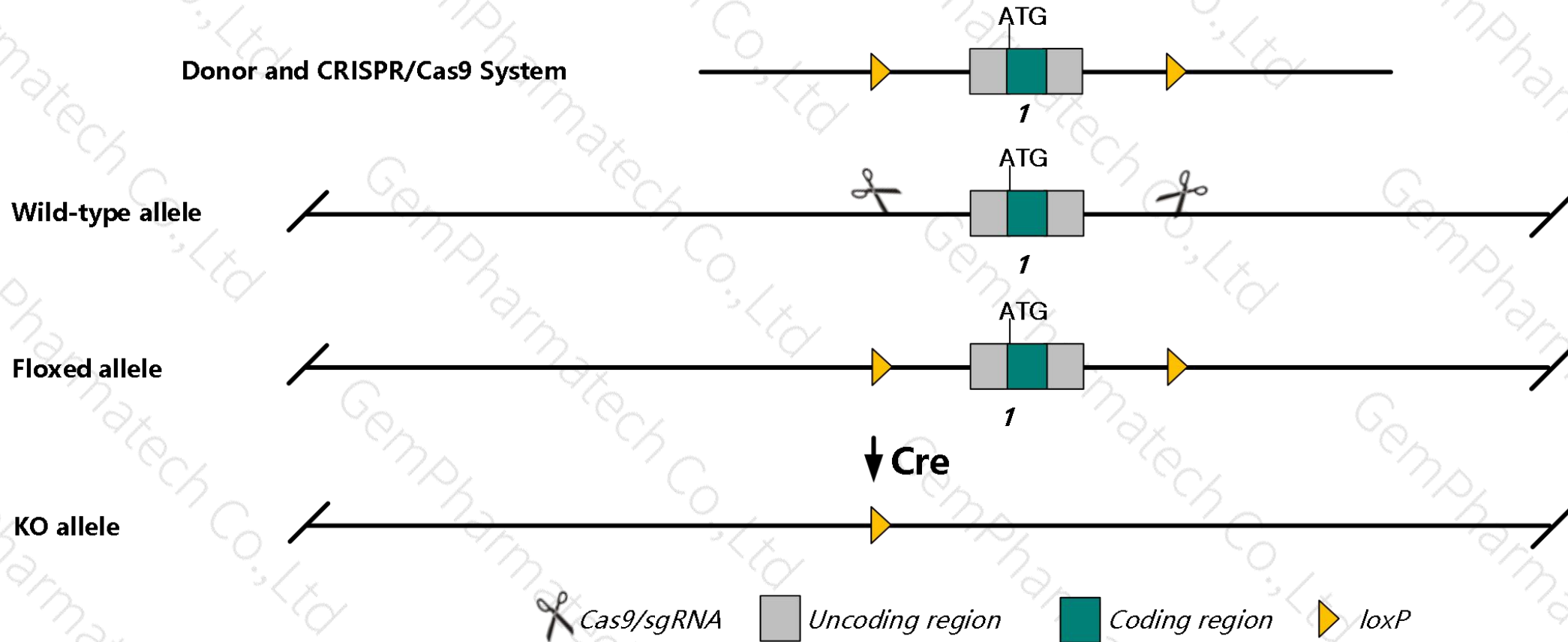
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gpr52* gene. The schematic diagram is as follows:



- The *Gpr52* gene has 1 transcript. According to the structure of *Gpr52* gene, exon1 of *Gpr52-201* (ENSMUST00000238289.1) 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 *Gpr52* 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.

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit decreased anxiety response in an open field test and increased startle response when treated with MK-801.
- *Gpr52* gene overlaps with the intron of *Rabgap11* gene, and the effect on *Rabgap11* gene is unknown.
- The *Gpr52* gene is located on the Chr1. 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)

## Gpr52 G protein-coupled receptor 52 [ *Mus musculus* (house mouse) ]

Gene ID: 620246, updated on 24-Oct-2019

### Summary



Official Symbol	Gpr52 provided by <a href="#">MGI</a>
Official Full Name	G protein-coupled receptor 52 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MG1:3643278</a>
See related	<a href="#">Ensembl:ENSMUSG00000118401</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	EG620246
Expression	Broad expression in cortex adult (RPKM 1.5), frontal lobe adult (RPKM 1.4) and 15 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

### Genomic context



Location: 1; 1 H2.1

Exon count: 2

See Gpr52 in [Genome Data Viewer](#)

# Transcript information (Ensembl)

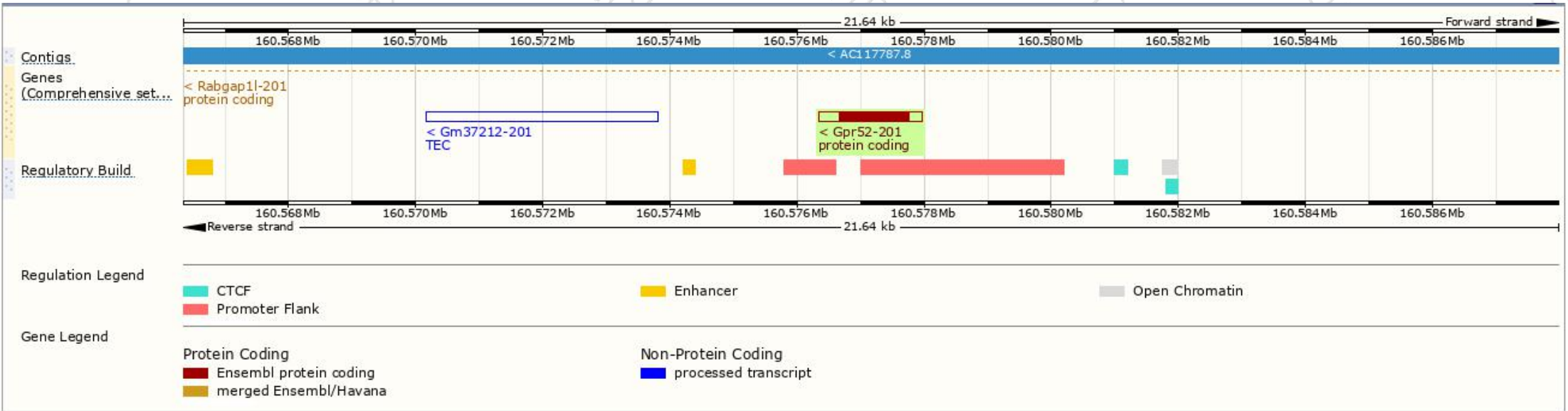
The gene has 1 transcript,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gpr52-201	<a href="#">ENSMUST00000238289.1</a>	1637	<a href="#">361aa</a>	<div><div></div>Protein coding</div>	-	<a href="#">P0C5J4</a>	GENCODE basic APPRIS P1

The strategy is based on the design of *Gpr52-201* transcript,The transcription is shown below

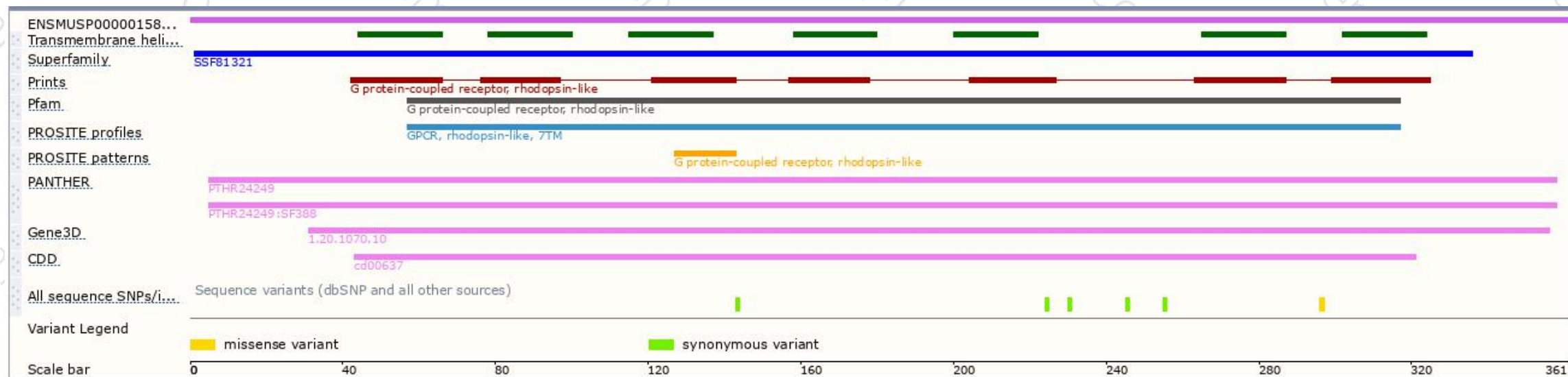


# Genomic location distribution

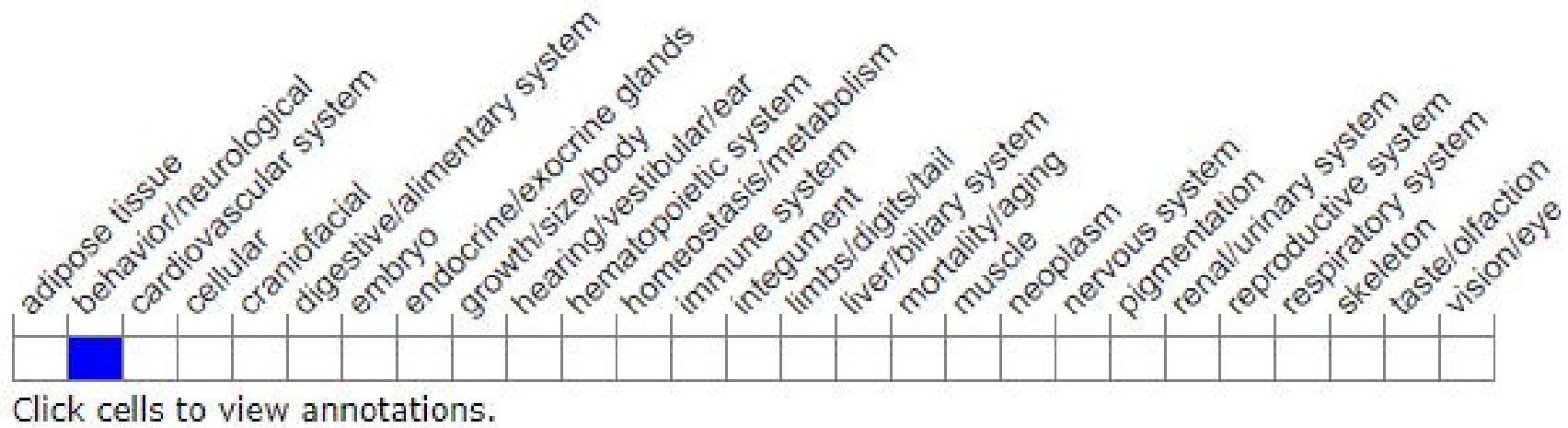




# Protein domain



# 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 knock-out allele exhibit decreased anxiety response in an open field test and increased startle response when treated with MK-801.

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

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