# Mrgprd Cas9-CKO Strategy

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## **Project Overview**



**Project Name** 

Mrgprd

**Project type** 

Cas9-CKO

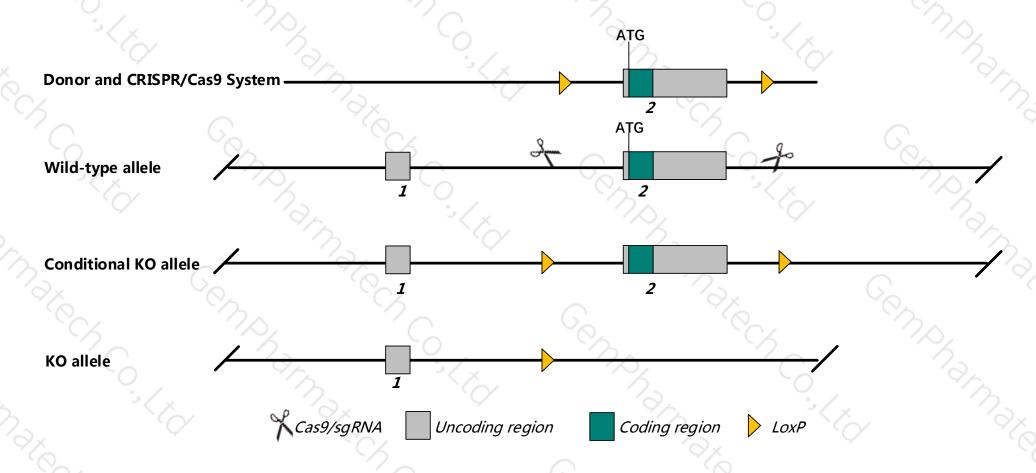
Strain background

C57BL/6JGpt

## **Conditional Knockout strategy**



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



### **Technical routes**



- ➤ The *Mrgprd* gene has 1 transcript.According to the structure of *Mrgprd* gene, exon2 of *Mrgprd*-201 (ENSMUST00000062163.7) transcript is recommended as the knockout region. The region contains all coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Mrgprd* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues or cell types.

### **Notice**



- According to the existing MGI data, mice homozygous for a knock-out allele exhibit decreased sensitivity to cold, heat, and mechanical stimuli. Mice for another null allele exhibit dilated cardiomypathy and cardiac dysfunction.
- The *Mrgprd* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

## Gene information (NCBI)



#### Mrgprd MAS-related GPR, member D [ Mus musculus (house mouse) ]

Gene ID: 211578, updated on 21-Aug-2019

#### Summary

☆ ?

Official Symbol Mrgprd provided by MGI

Official Full Name MAS-related GPR, member D provided by MGI

Primary source MGI:MGI:3033142

See related Ensembl: ENSMUSG00000051207

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 Mgrd; MrgD; TGR7; Gm499

Expression Low expression observed in reference dataset See more

Orthologs human all

# Transcript information (Ensembl)



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

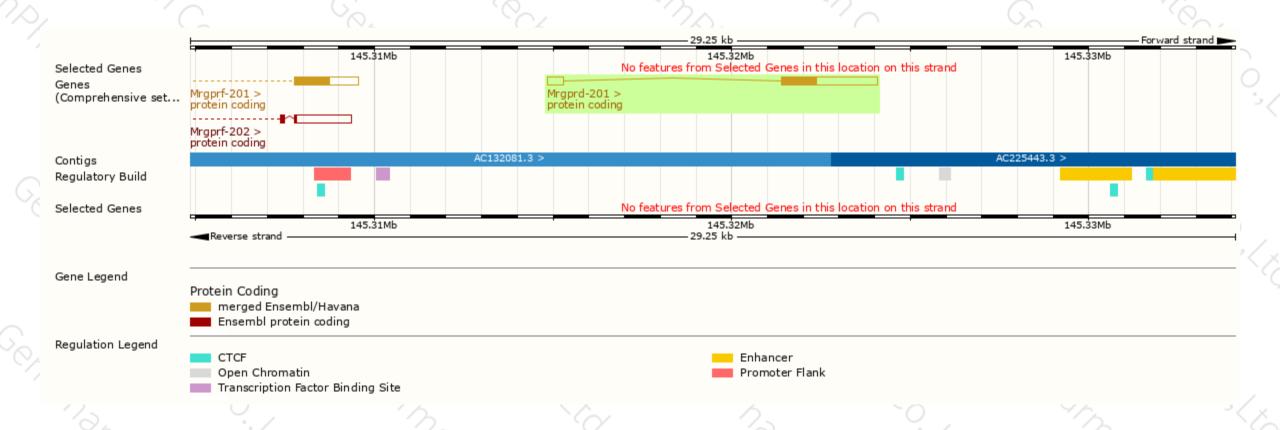
Name 🍦	Transcript ID	bp 🛊	Protein	Biotype	CCDS	UniProt ▼		Flags	\$
Mrgprd-201	ENSMUST00000062163.7	3144	<u>321aa</u>	Protein coding	CCDS22057₺	<u>Q5UCB4</u> & <u>Q91ZB8</u> &	TSL:1	GENCODE basic	APPRIS P1

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



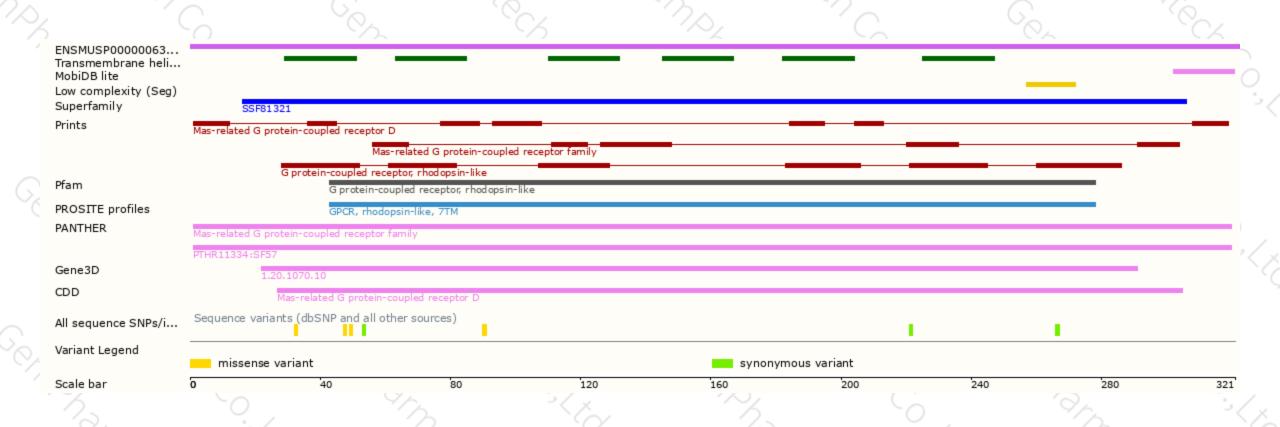
### Genomic location distribution





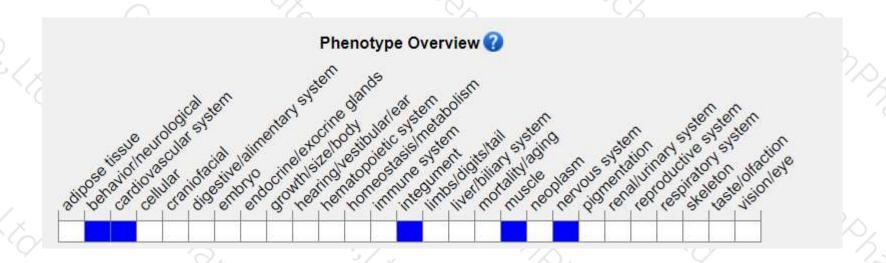
### Protein domain





## Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

Mice homozygous for a knock-out allele exhibit decreased sensitivity to cold, heat, and mechanical stimuli. Mice for another null allele exhibit dilated cardiomypathy and cardiac dysfunction.

If you have any questions, you are welcome to inquire. Tel: 025-5864 1534





