

# Prss37 Cas9-CKO Strategy

Designer: Daohua Xu

**Design Date:** 2019-7-30

# **Project Overview**



**Project Name** 

Prss37

**Project type** 

Cas9-CKO

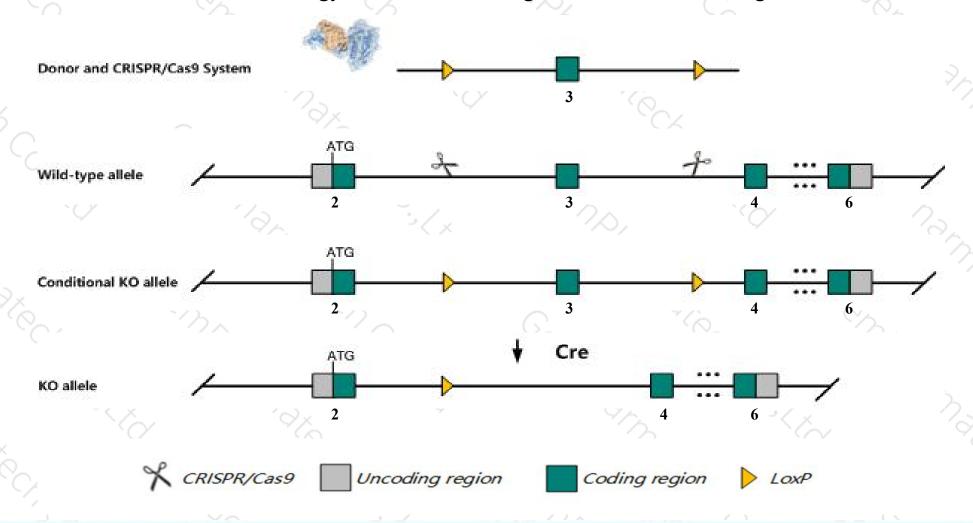
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Prss37* gene. The schematic diagram is as follows:



### Technical routes



- ➤ The *Prss37* gene has 2 transcripts. According to the structure of *Prss37* gene, exon3 of *Prss37-201*(ENSMUST00000031967.5) transcript is recommended as the knockout region. The region contains 142bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Prss37* 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 knock-out allele exhibit male infertility associated with impaired zona binding and migration from the uterus into the oviduct.
- The *Prss37* gene is located on the Chr6. 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)



#### Prss37 protease, serine 37 [Mus musculus (house mouse)]

Gene ID: 67690, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Prss37 provided by MGI

Official Full Name protease, serine 37 provided by MGI

Primary source MGI:MGI:1914940

See related Ensembl:ENSMUSG00000029909

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 1700016G05Rik, Tryx2

Expression Restricted expression toward testis adult (RPKM 93.1)See more

Orthologs <u>human</u> all

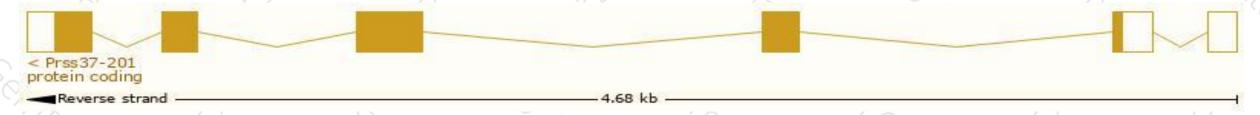
# Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

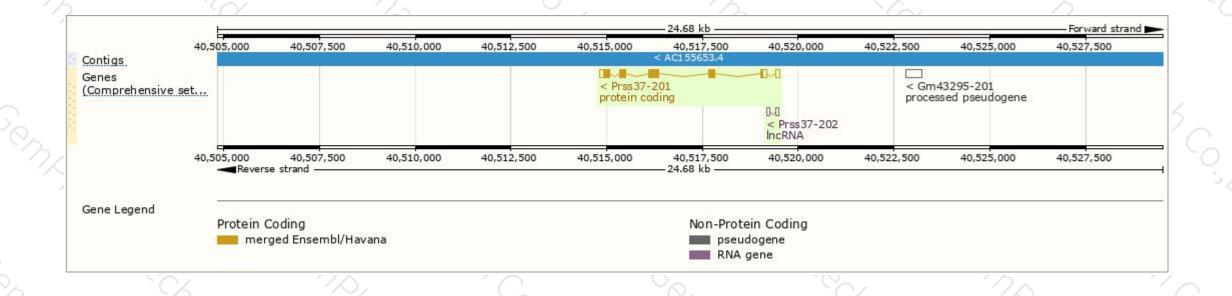
Name 🍦	Transcript ID	bp 🌲	Protein 🍦	Biotype 👙	CCDS 🍦	UniProt	Flags		
Prss37-201	ENSMUST00000031967.5	1049	<u>237aa</u>	Protein coding	CCDS20033₫	Q9DAA4₫	TSL:1	GENCODE basic	APPRIS P1
Prss37-202	ENSMUST00000148195.1	175	No protein	IncRNA	100	(C <del>t.</del> )		TSL:5	

The strategy is based on the design of *Prss37-201* transcript, The transcription is shown below



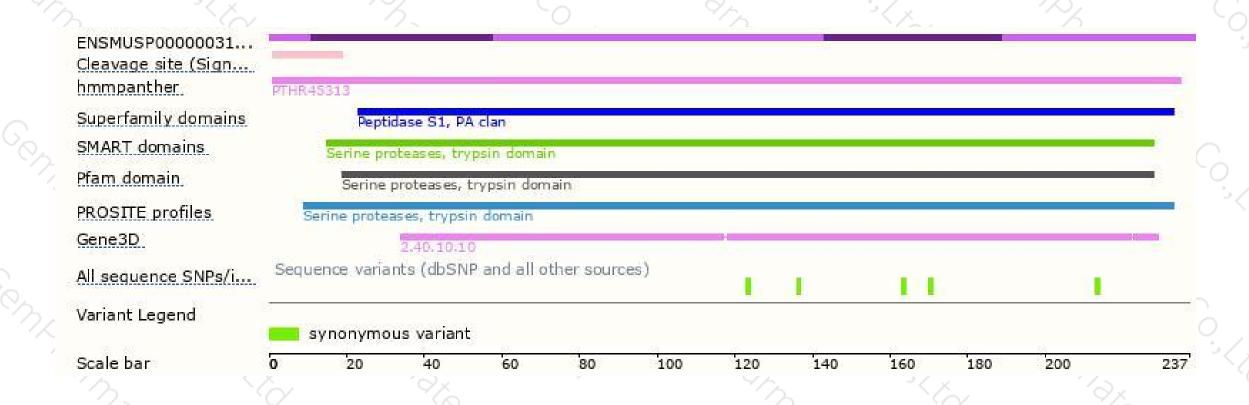
### Genomic location distribution





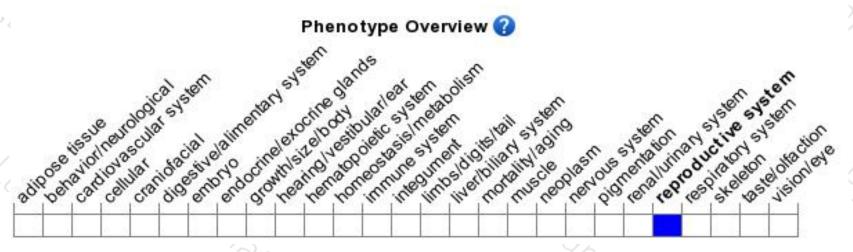
### Protein domain





# 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 male infertility associated with impaired zona binding and migration from the uterus into the oviduct.



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





