

Prss37 Cas9-KO Strategy

Designer:

Daohua Xu

Design Date:

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Project Overview

Project Name

Prss37

Project type

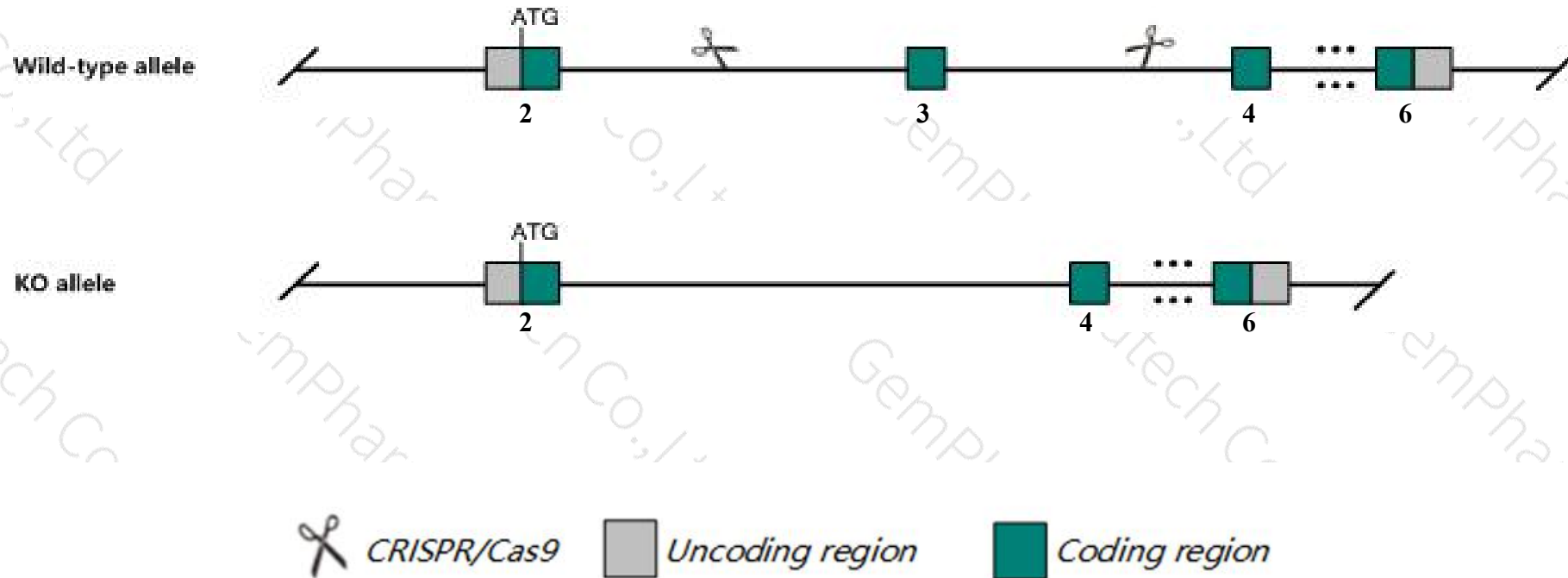
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- 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

- 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

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

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

Summary



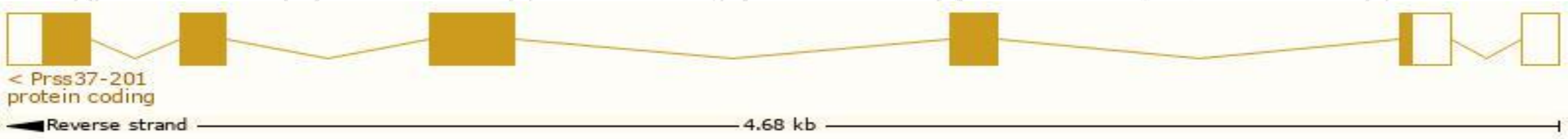
Official Symbol	Prss37 provided by MGI
Official Full Name	protease, serine 37 provided by MGI
Primary source	MGI:MGI:1914940
See related	Ensembl:ENSMUSG00000029909
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; 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	human all

Transcript information (Ensembl)

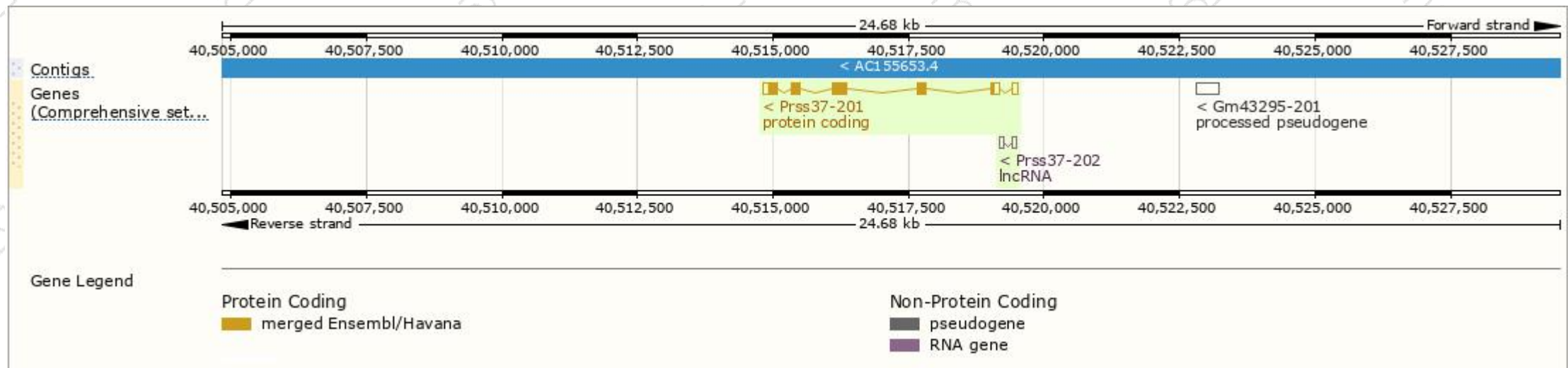
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Prss37-201	ENSMUST00000031967.5	1049	237aa	Protein coding	CCDS20033	Q9DAA4	TSL:1 GENCODE basic APPRIS P1
Prss37-202	ENSMUST00000148195.1	175	No protein	lncRNA	-	-	TSL:5

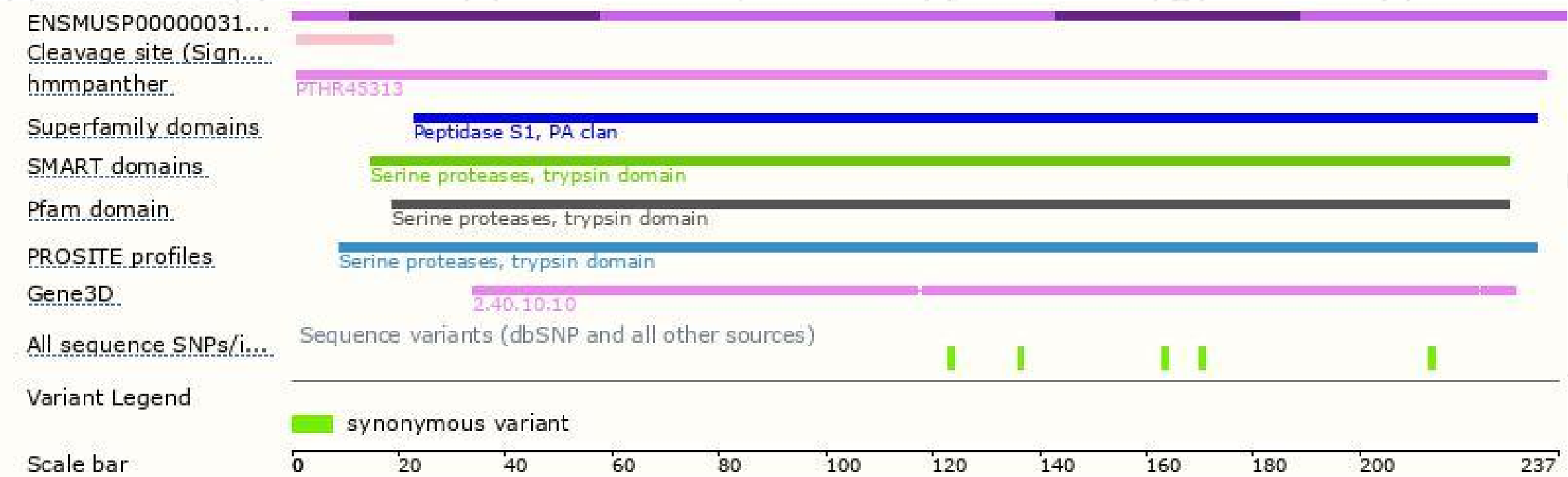
The strategy is based on the design of *Prss37-201* transcript,The transcription is shown below



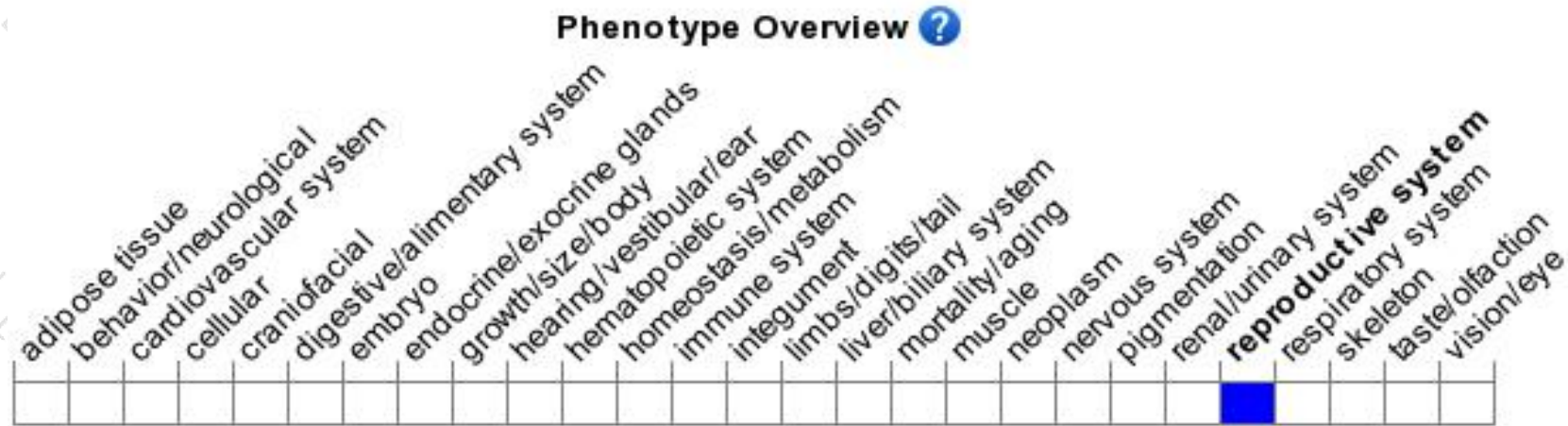
Genomic location distribution



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

