

Prss50 Cas9-CKO Strategy

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Project Overview

Project Name

Prss50

Project type

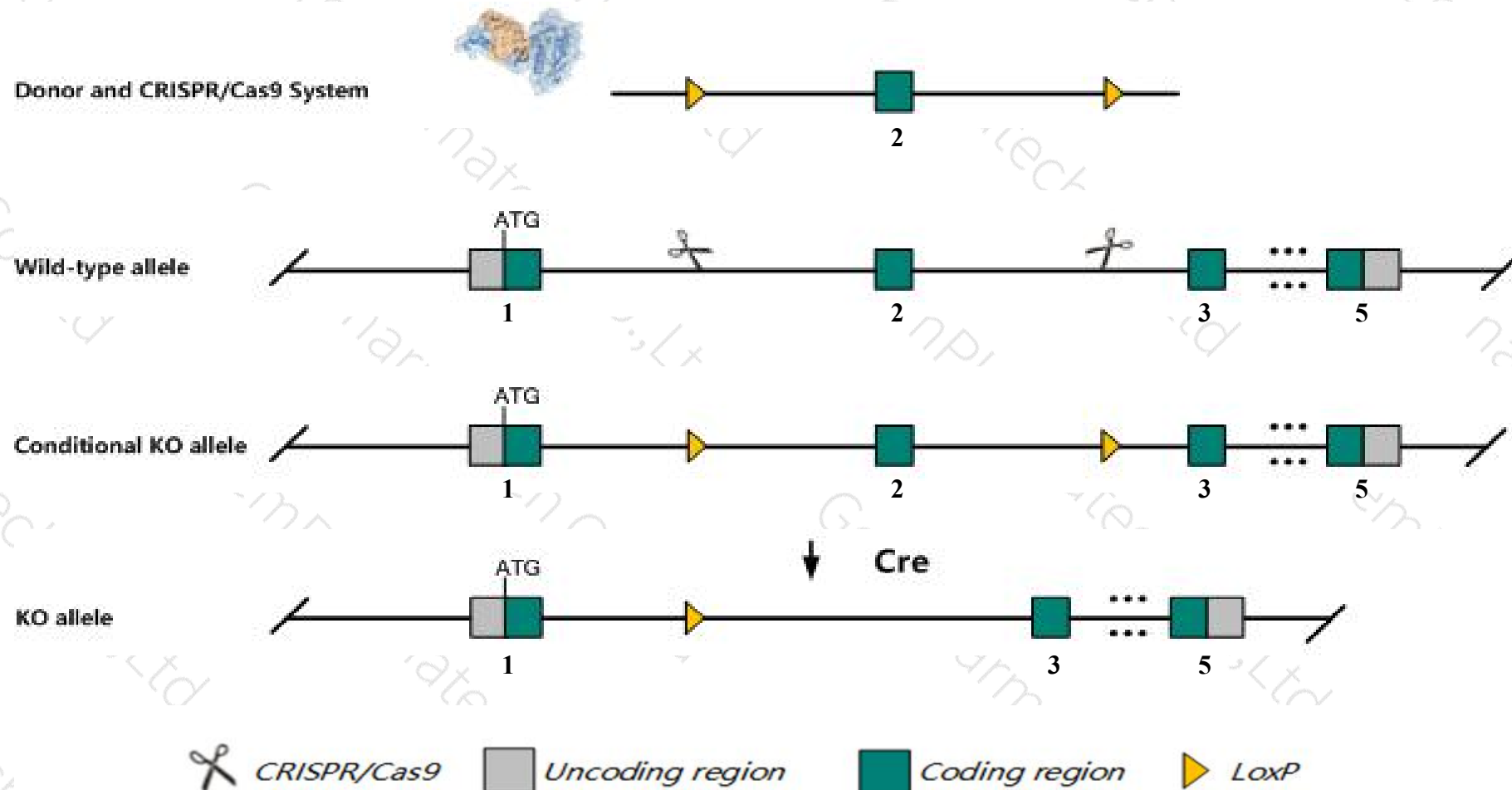
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Prss50* gene. The schematic diagram is as follows:



- The *Prss50* gene has 2 transcripts. According to the structure of *Prss50* gene, exon2 of *Prss50-201* (ENSMUST00000051097.5) transcript is recommended as the knockout region. The region contains 163bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Prss50* 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

- The *Prss50* gene is located on the Chr9. 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)

Prss50 protease, serine 50 [Mus musculus (house mouse)]

Gene ID: 235631, updated on 31-Jan-2019

Summary



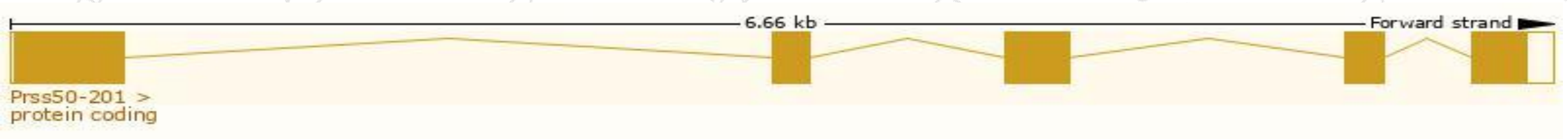
Official Symbol	Prss50 provided by MGI
Official Full Name	protease, serine 50 provided by MGI
Primary source	MGI:MGI:2447303
See related	Ensembl:ENSMUSG00000048752
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	79H19C, Tsp50
Expression	Biased expression in testis adult (RPKM 29.0), liver E14.5 (RPKM 5.0) and 2 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

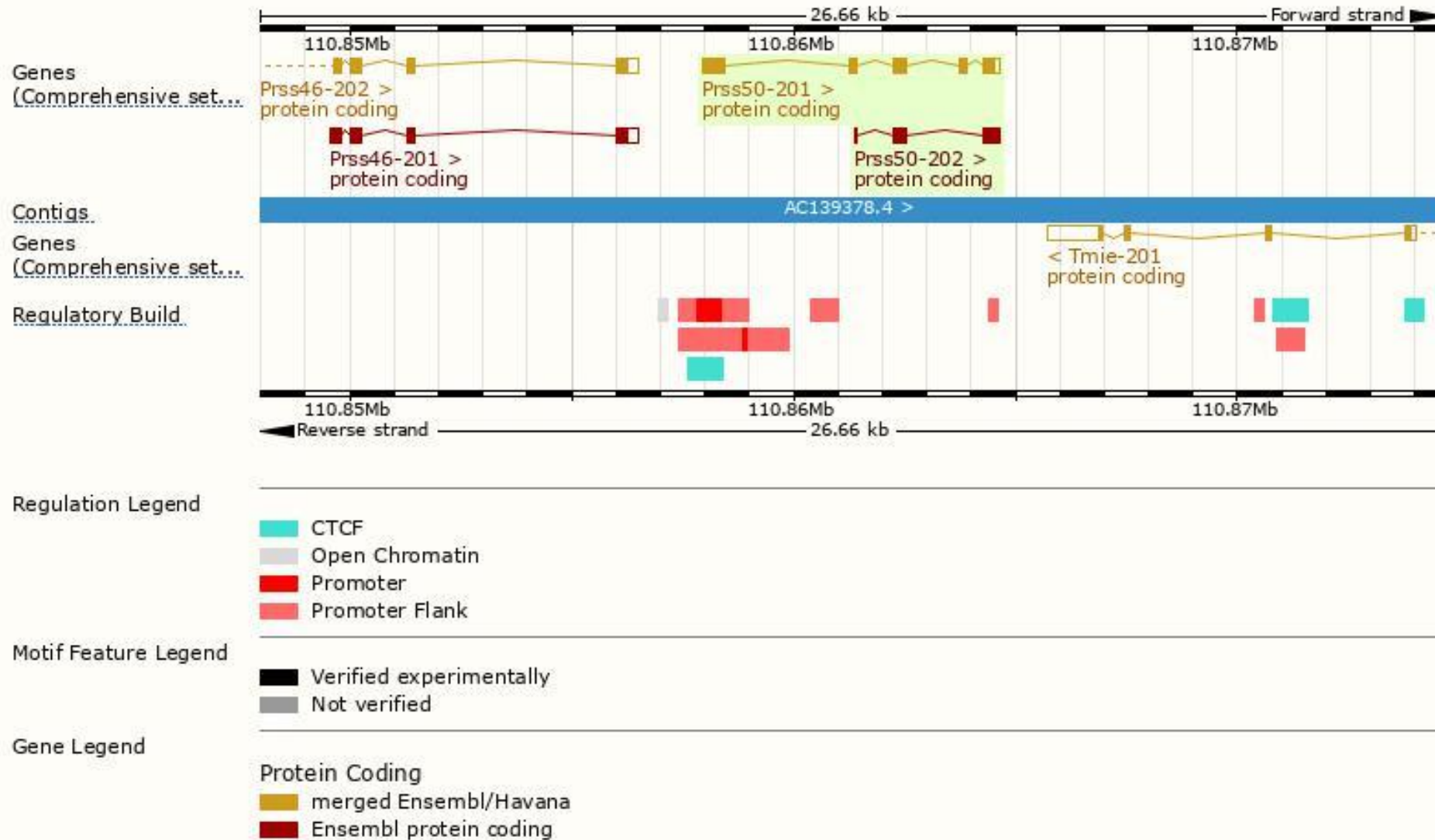
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Prss50-201	ENSMUST00000051097.5	1456	439aa	Protein coding	CCDS23577	Q8BLH5	TSL:1 GENCODE basic APPRIS P1
Prss50-202	ENSMUST00000196027.1	689	221aa	Protein coding	-	A0A0G2JFA6	CDS 5' incomplete TSL:3

The strategy is based on the design of *Prss50-201* transcript,The transcription is shown below



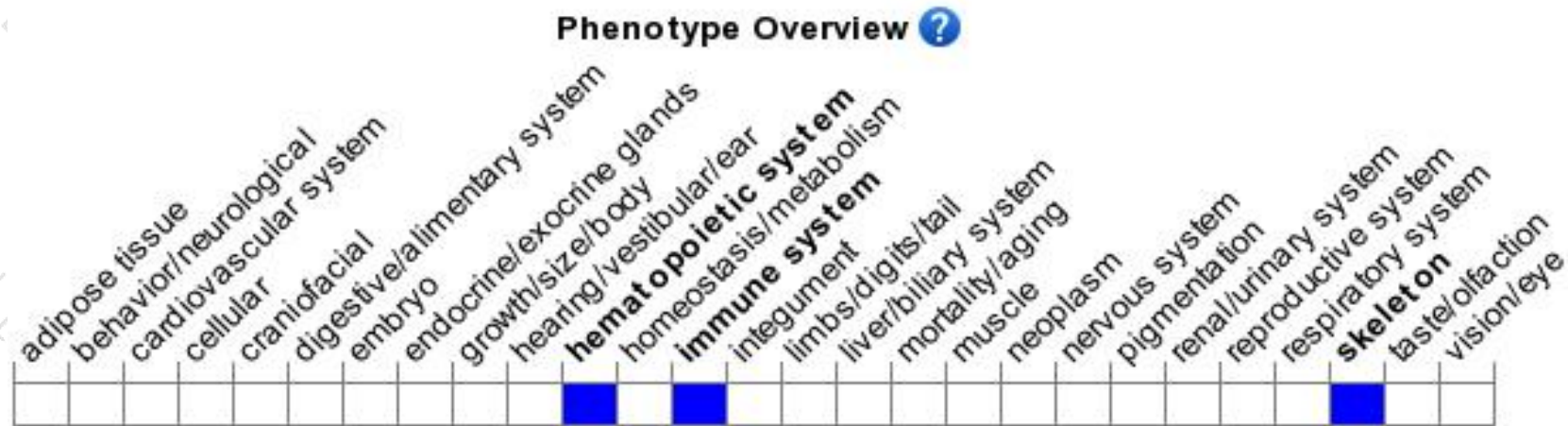
Genomic location distribution



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/>).

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

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