

Prss22 Cas9-KO Strategy

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Project Overview

Project Name

Prss22

Project type

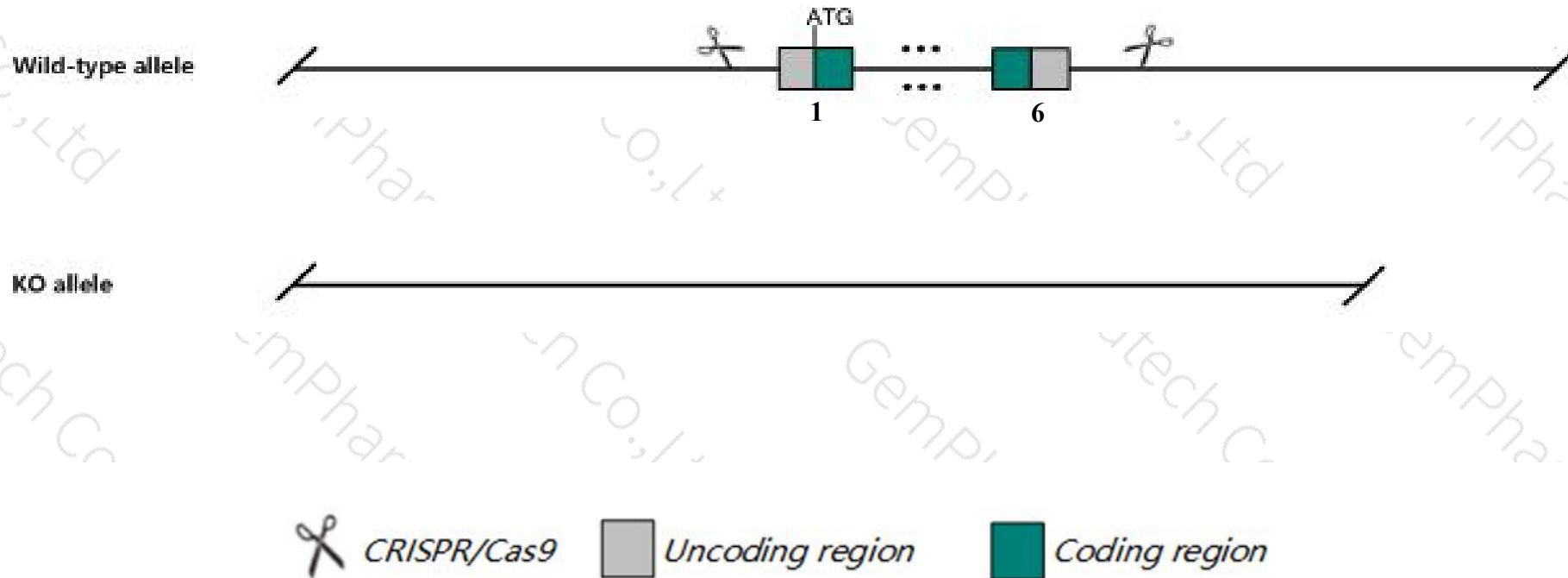
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Prss22* gene. The schematic diagram is as follows:



- The *Prss22* gene has 1 transcript. According to the structure of *Prss22* gene, exon1-exon6 of *Prss22-201* (ENSMUST00000041649.7) 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 *Prss22* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Prss22* gene is located on the Chr17. 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)

Prss22 protease, serine 22 [Mus musculus (house mouse)]

Gene ID: 70835, updated on 15-Feb-2019

Summary



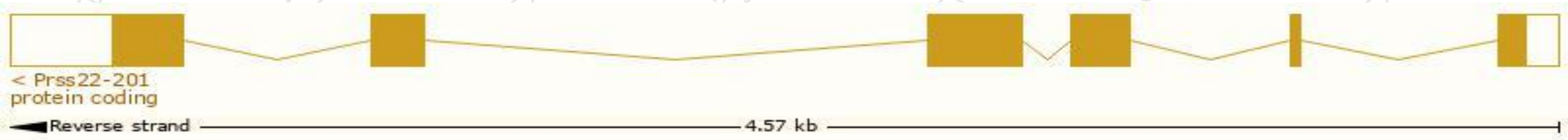
Official Symbol	Prss22 provided by MGI
Official Full Name	protease, serine 22 provided by MGI
Primary source	MGI:MGI:1918085
See related	Ensembl:ENSMUSG00000045027
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4733401N09Rik, BSSP-4, SP001LA
Expression	Biased expression in bladder adult (RPKM 6.1), stomach adult (RPKM 0.6) and 4 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

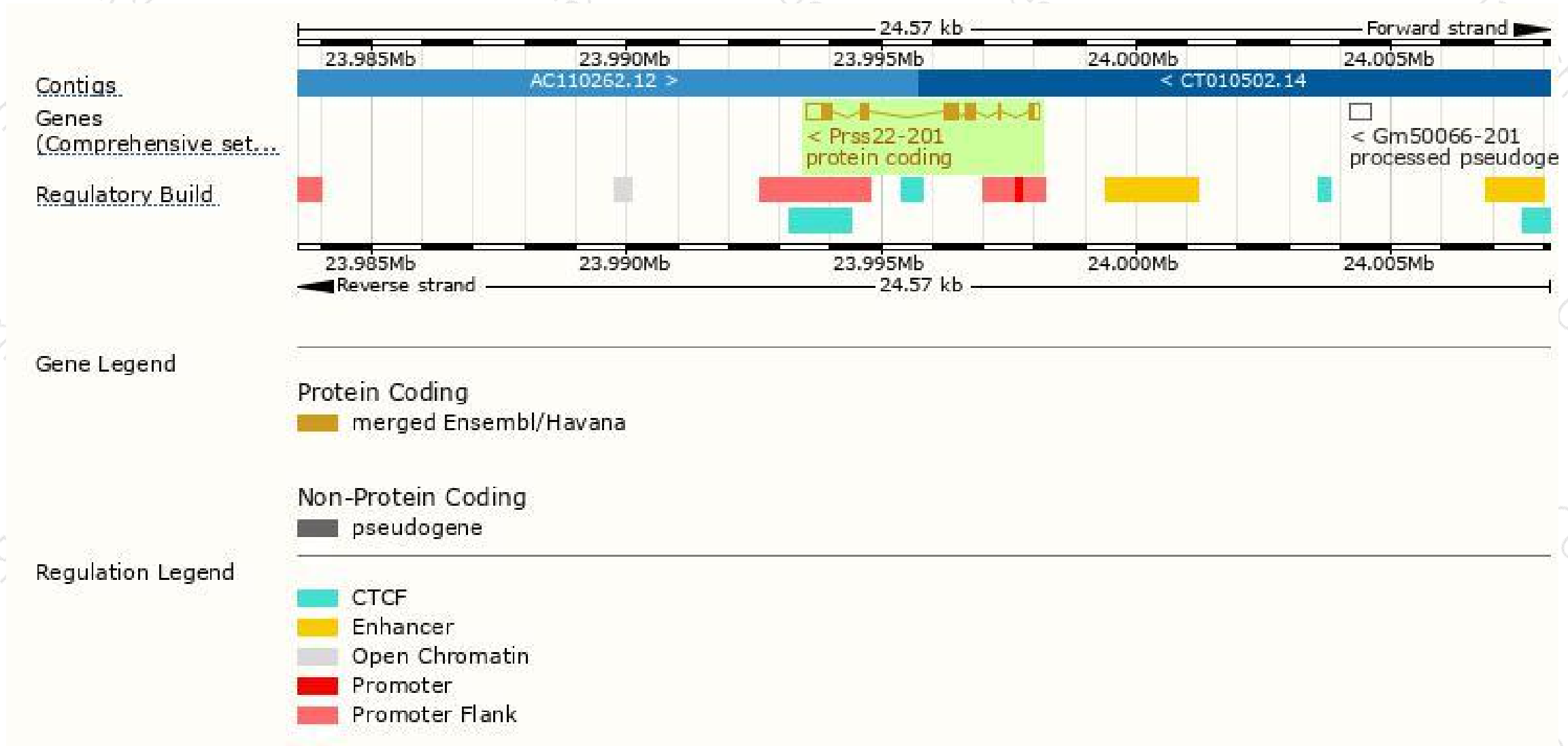
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Prss22-201	ENSMUST00000041649.7	1324	307aa	Protein coding	CCDS37482	Q7TML0	TSL:1 GENCODE basic APPRIS P1

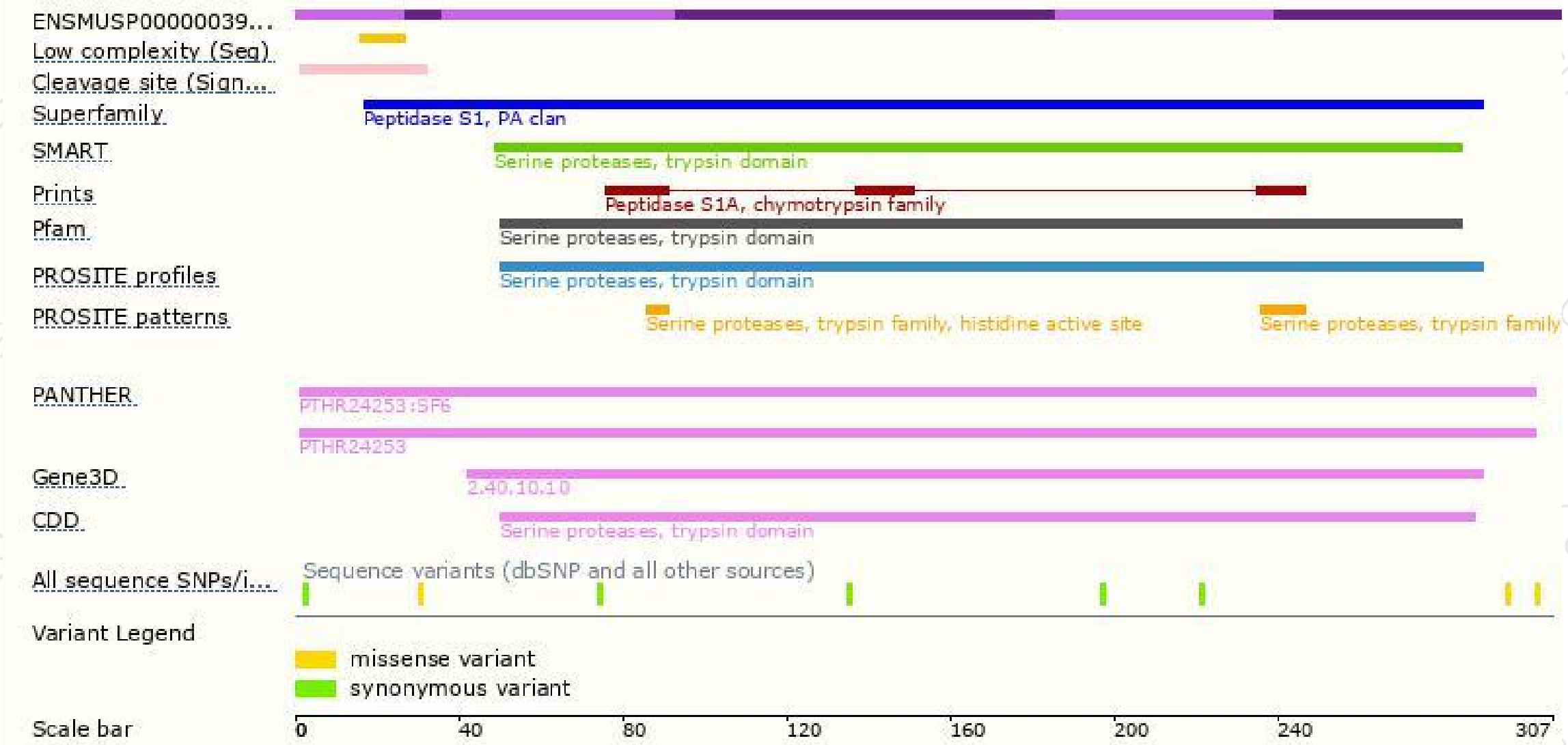
The strategy is based on the design of *Prss22-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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