

Pros1 Cas9-CKO Strategy

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Project Overview



Project Name

Pros1

Project type

Cas9-CKO

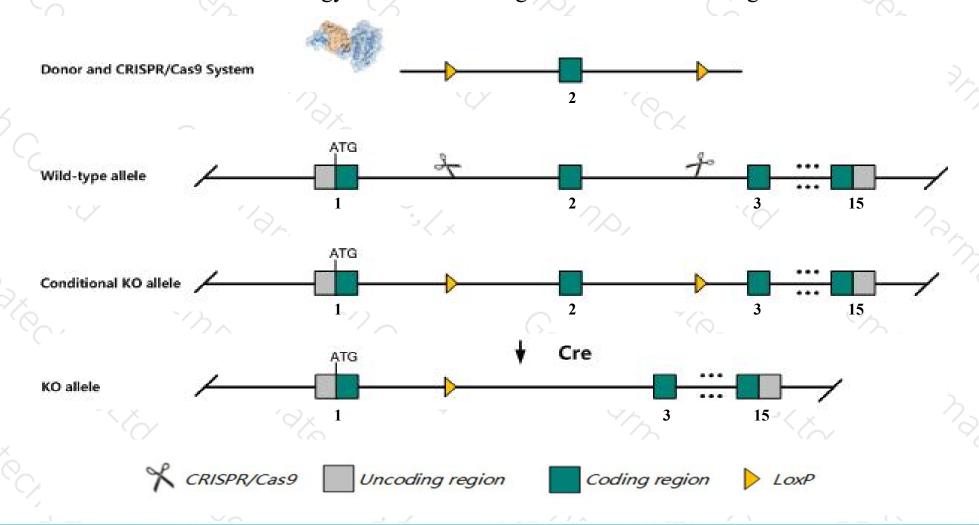
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pros1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Pros1* gene has 3 transcripts. According to the structure of *Pros1* gene, exon2 of *Pros1-201*(ENSMUST00000023629.8) transcript is recommended as the knockout region. The region contains 158bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Pros1* 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 neonatal lethality associated with thrombosis, hemorrhage, and thrombocytopenia.
- ➤ Transcript 202, 203 are unaffected.
- The *Pros1* gene is located on the Chr16. 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)



Pros1 protein S (alpha) [Mus musculus (house mouse)]

Gene ID: 19128, updated on 12-Aug-2019

Summary

△ ?

Official Symbol Pros1 provided by MGI

Official Full Name protein S (alpha) provided by MGI

Primary source MGI:MGI:1095733

See related Ensembl: ENSMUSG00000022912

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as AW214361

Summary This gene encodes a vitamin K-dependent protein with key roles in multiple biological processes including coagulation, apoptosis and

vasculogenesis. The encoded protein undergoes proteolytic processing to generate a mature protein which is secreted into the plasma.

Mice lacking the encoded protein die in utero from a fulminant coagulopathy and associated hemorrhages. [provided by RefSeq. Oct

2015]

Expression Ubiquitous expression in placenta adult (RPKM 16.0), bladder adult (RPKM 11.9) and 27 other tissues See more

Orthologs human all

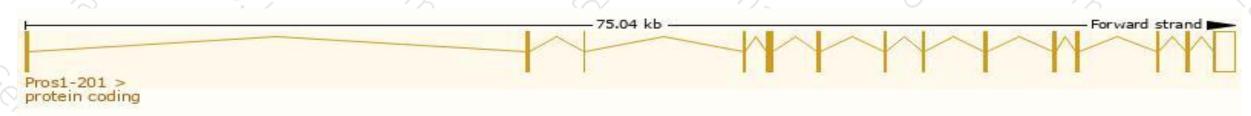
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

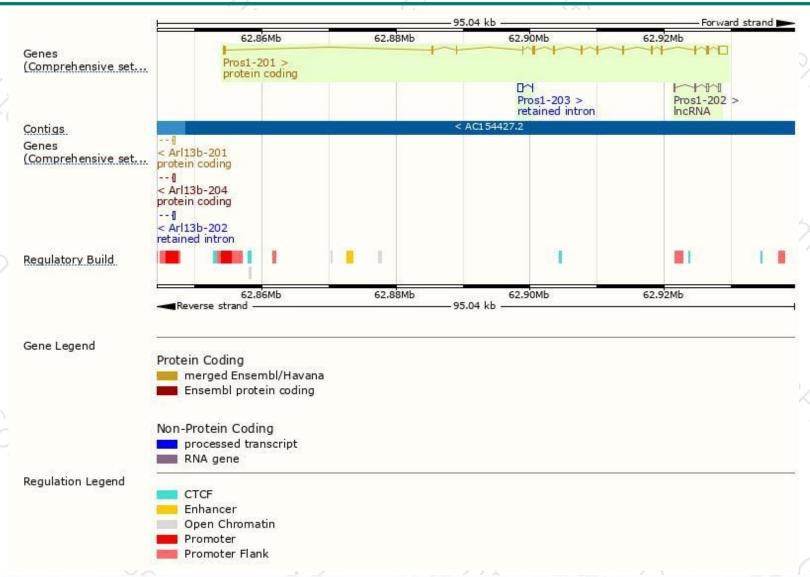
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pros1-201	ENSMUST00000023629.8	3303	<u>675aa</u>	Protein coding	CCDS28263	Q08761 Q3TR66	TSL:1 GENCODE basic APPRIS P1
Pros1-203	ENSMUST00000155940.1	863	No protein	Retained intron		*	TSL:5
Pros1-202	ENSMUST00000127502.1	791	No protein	IncRNA	ÿ <u>4</u>	-	TSL:1

The strategy is based on the design of *Pros1-201* transcript, The transcription is shown below



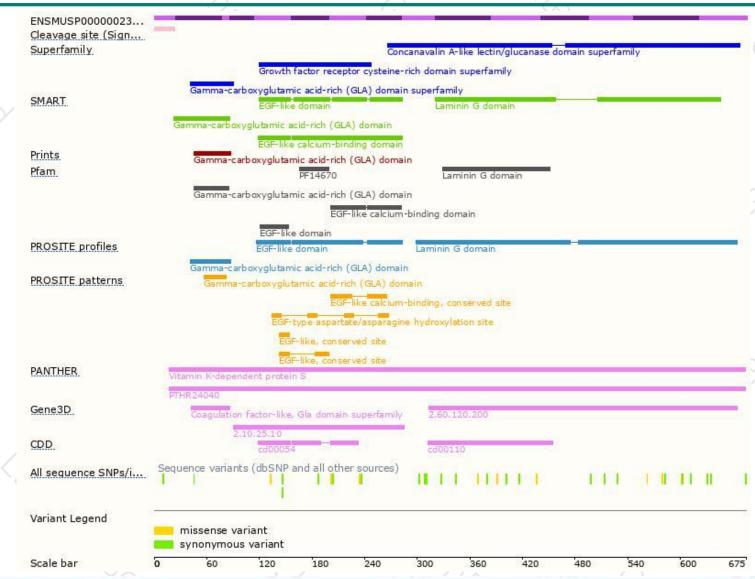
Genomic location distribution





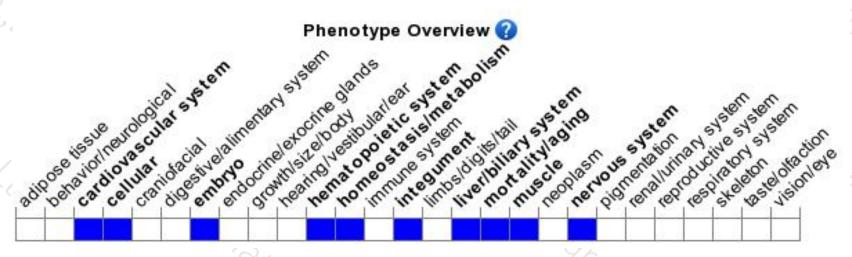
Protein domain





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 neonatal lethality associated with thrombosis, hemorrhage, and thrombocytopenia.



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





