

# Proc Cas9-KO Strategy

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### **Project Overview**



Project Name Proc

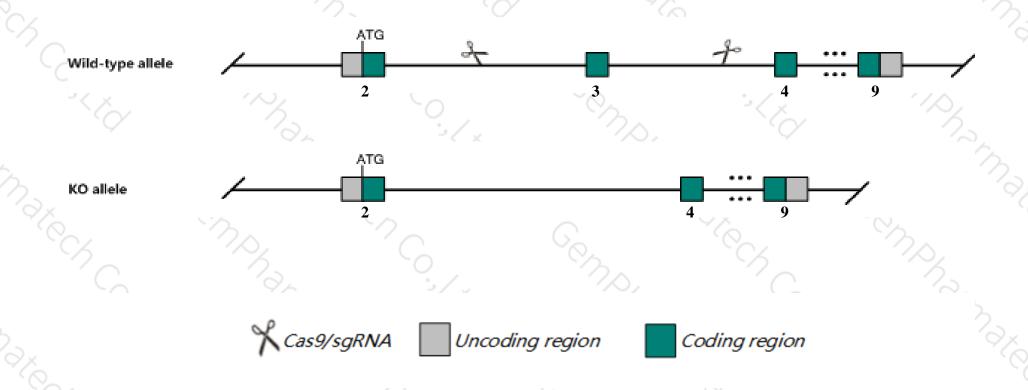
Project type Cas9-KO

Strain background C57BL/6JGpt

## **Knockout strategy**



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



### **Technical routes**



- ➤ The *Proc* gene has 8 transcripts. According to the structure of *Proc* gene, exon3 of *Proc-201*(ENSMUST00000171765.1) transcript is recommended as the knockout region. The region contains 164bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Proc* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA 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.

### **Notice**



- ➤ According to the existing MGI data, Inactivation of the locus results in death within 24 hours of birth due to consumptive coagulopathy. Thromboses and bleeding are observed in the brains and livers of homozygous mutant mice.
- ➤ The *Proc* gene is located on the Chr18. 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)



#### Proc protein C [ Mus musculus (house mouse) ]

Gene ID: 19123, updated on 12-Aug-2019

Summary

☆ ?

Official Symbol Proc provided by MGI

Official Full Name protein C provided by MGI

Primary source MGI:MGI:97771

See related Ensembl: ENSMUSG00000024386

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 PC

Summary This gene encodes the vitamin K-dependent protein C, which plays a vital role in the anticoagulation pathway. The encoded protein undergoes proteolytic processing

including activation by thrombin-thrombomodulin complex to form the anticoagulant serine protease that degrades activated coagulation factors. A complete lack of the encoded protein in mice results in severe perinatal consumptive coagulopathy in the brain and liver, resulting in death within 24 hours after birth. Alternative splicing results in multiple transcript variants encoding different isoforms that may undergo similar processing to generate the mature protein. [provided by RefSeq.

Sep 2015]

Expression Biased expression in liver adult (RPKM 230.9), liver E18 (RPKM 159.9) and 4 other tissues See more

Orthologs human all

# Transcript information (Ensembl)



The gene has 8 transcripts, all transcripts are shown below:

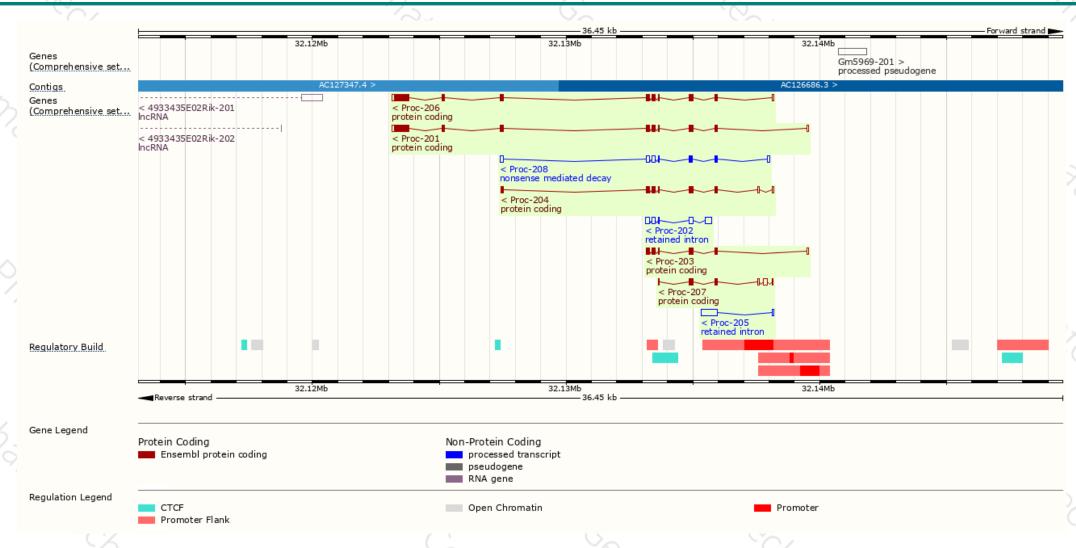
>	Name 🌲	Transcript ID	bp 🌲	Protein	Biotype	CCDS 🍦	UniProt 🍦	Flags ♦
1	Proc-206	ENSMUST00000234651.1	1575	<u>460aa</u>	Protein coding	CCDS29116₽	-	GENCODE basic   APPRIS P1
	Proc-201	ENSMUST00000171765.1	1566	<u>460aa</u>	Protein coding	CCDS29116₽	<u>P33587</u> ₽	TSL:5 GENCODE basic APPRIS P1
	Proc-204	ENSMUST00000234386.1	781	<u>214aa</u>	Protein coding	-	-	CDS 3' incomplete
	Proc-203	ENSMUST00000234375.1	614	<u>177aa</u>	Protein coding	-	-	CDS 3' incomplete
	Proc-207	ENSMUST00000234657.1	558	<u>86aa</u>	Protein coding	-	-	CDS 3' incomplete
	Proc-208	ENSMUST00000234746.1	719	<u>51aa</u>	Nonsense mediated decay	-	-	-
	Proc-202	ENSMUST00000234198.1	733	No protein	Retained intron	-	-	-
	Proc-205	ENSMUST00000234649.1	659	No protein	Retained intron	-	-	-

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



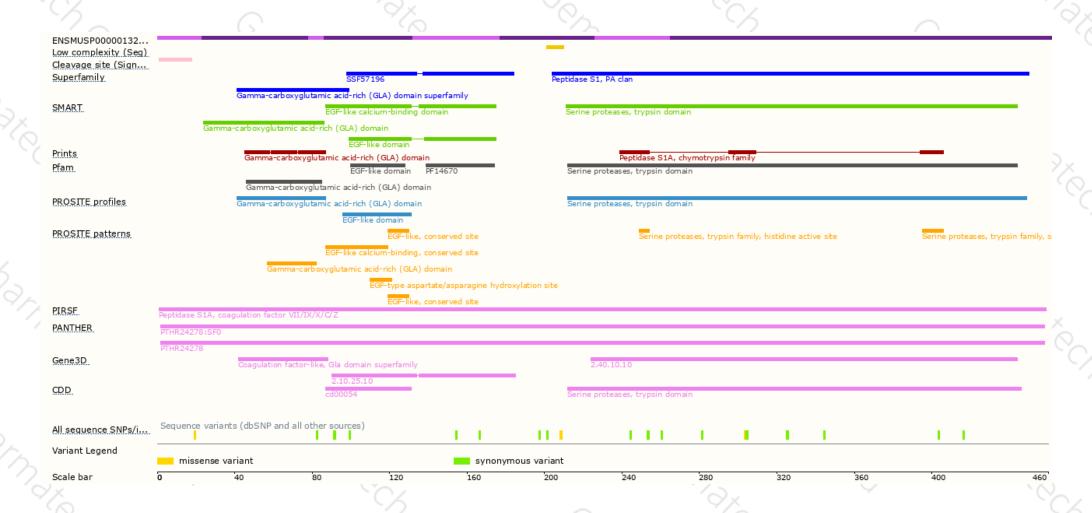
### Genomic location distribution





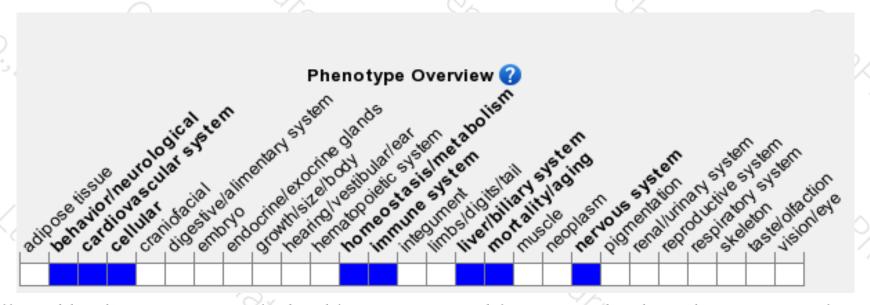
### Protein domain





### Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(http://www.informatics.jax.org/).

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

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