

Api5 Cas9-CKO Strategy

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Project Overview



Project Name

Api5

Project type

Cas9-CKO

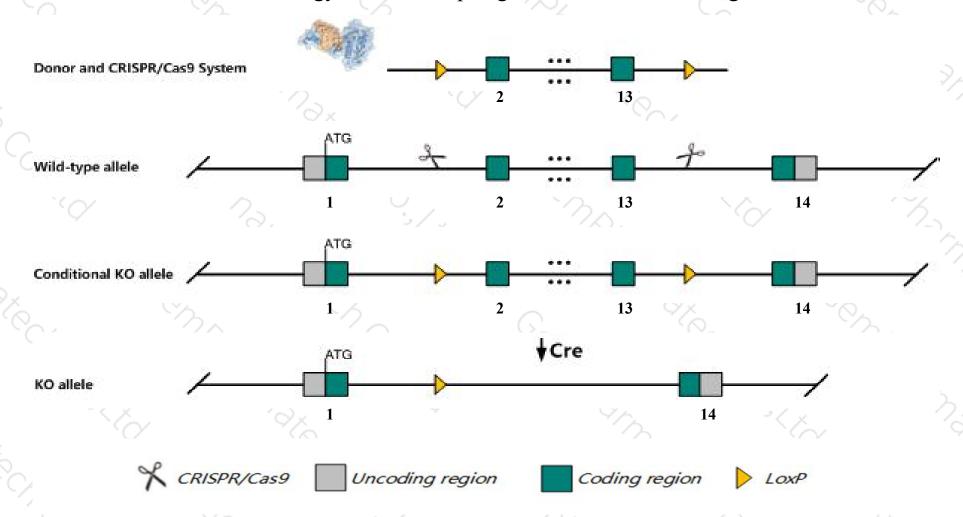
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Api5 gene. The schematic diagram is as follows:



Technical routes



- ➤ The Api5 gene has 4 transcripts. According to the structure of Api5 gene, exon2-exon13 of Api5-201

 (ENSMUST00000028617.6) transcript is recommended as the knockout region. The region contains 1423bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Api5* 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 *Api5* gene is located on the Chr2. 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)



Api5 apoptosis inhibitor 5 [Mus musculus (house mouse)]

Gene ID: 11800, updated on 4-May-2020

Summary

↑ ?

Official Symbol Api5 provided by MGI

Official Full Name apoptosis inhibitor 5 provided by MGI

Primary source MGI:MGI:1888993

See related Ensembl: ENSMUSG00000027193

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 API-5; AAC-11; AI196452

Expression Ubiquitous expression in CNS E11.5 (RPKM 28.3), bladder adult (RPKM 22.1) and 28 other tissues See more

Orthologs <u>human</u> all

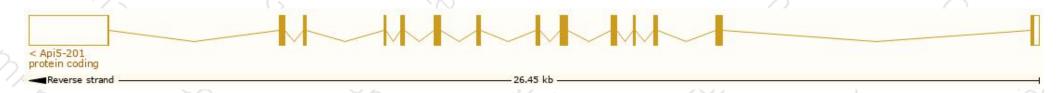
Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

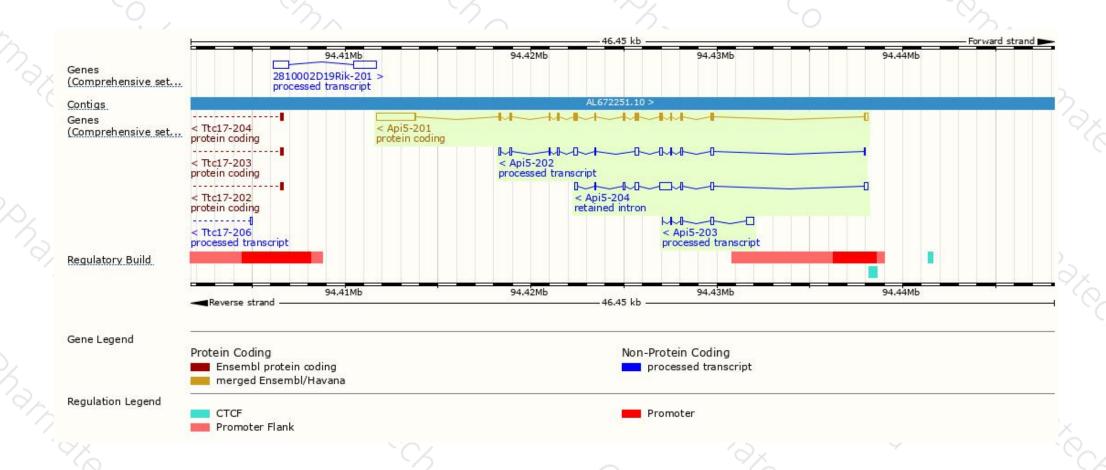
Name 🍦	Transcript ID	bp 🛊	Protein	Biotype	CCDS	UniProt	Flags
Api5-201	ENSMUST00000028617.6	3736	<u>504aa</u>	Protein coding	CCDS16459 ₽	<u>O35841</u> €	TSL:1 GENCODE basic APPRIS P1
Api5-202	ENSMUST00000144390.7	1308	No protein	Processed transcript	-	('	TSL:5
Api5-203	ENSMUST00000150930.1	730	No protein	Processed transcript	-	(+)	TSL:3
Api5-204	ENSMUST00000152454.7	1659	No protein	Retained intron		14	TSL:2

The strategy is based on the design of Api5-201 transcript, the transcription is shown below



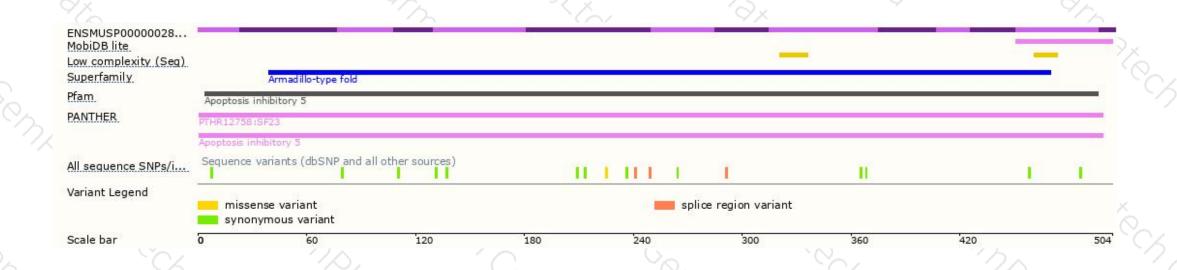
Genomic location distribution





Protein domain







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





