

Api5 Cas9-KO Strategy

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Project Overview

Project Name

Api5

Project type

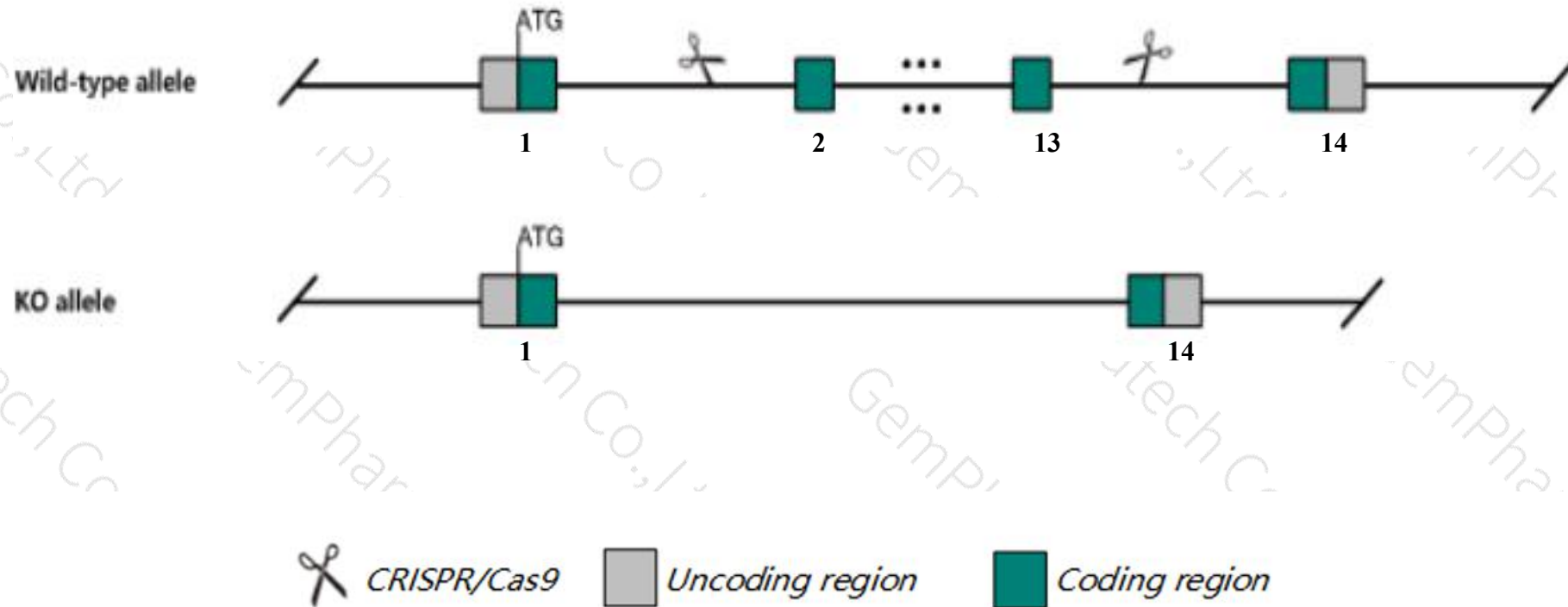
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- 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 was

- 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

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

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

Summary

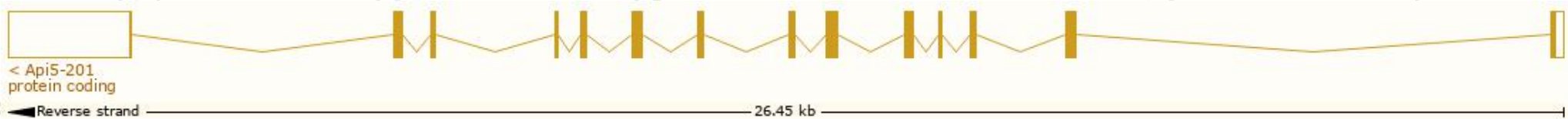
Official Symbol	Api5 provided by MGI
Official Full Name	apoptosis inhibitor 5 provided by MGI
Primary source	MGI:MGI:1888993
See related	Ensembl:ENSMUSG00000027193
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; 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	human all

Transcript information (Ensembl)

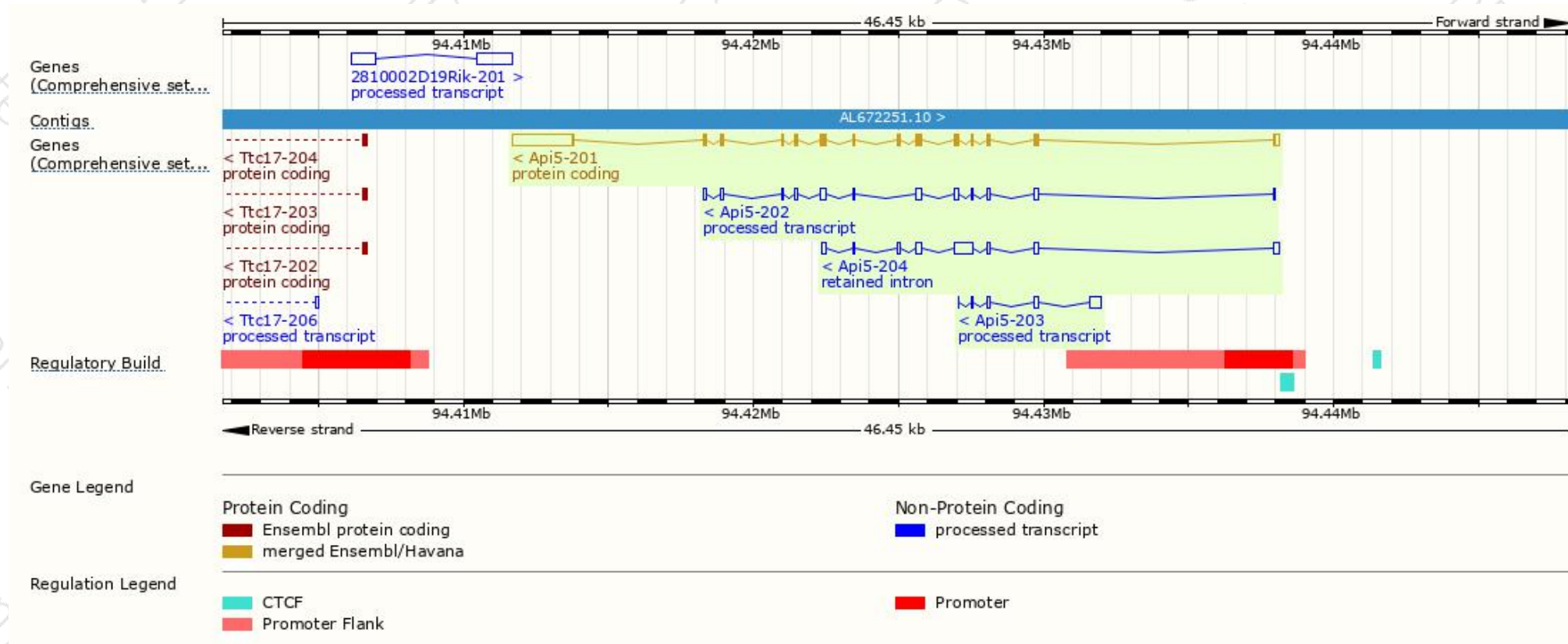
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Api5-201	ENSMUST00000028617.6	3736	504aa	Protein coding	CCDS16459	O35841	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	-	-	TSL:2

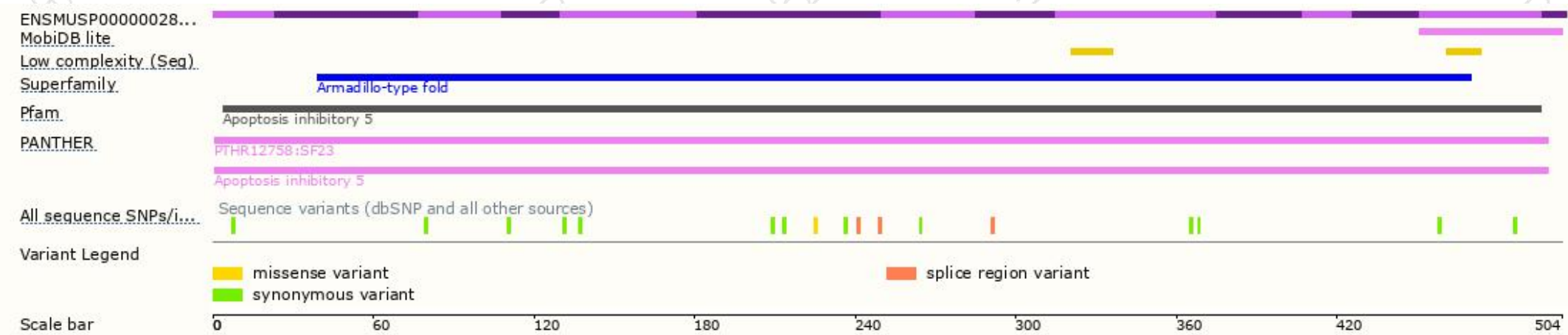
The strategy is based on the design of *Api5-201* transcript, the transcription is shown below



Genomic location distribution



Protein domain



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

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