

Capn7 Cas9-KO Strategy

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Project Overview

Project Name

Capn7

Project type

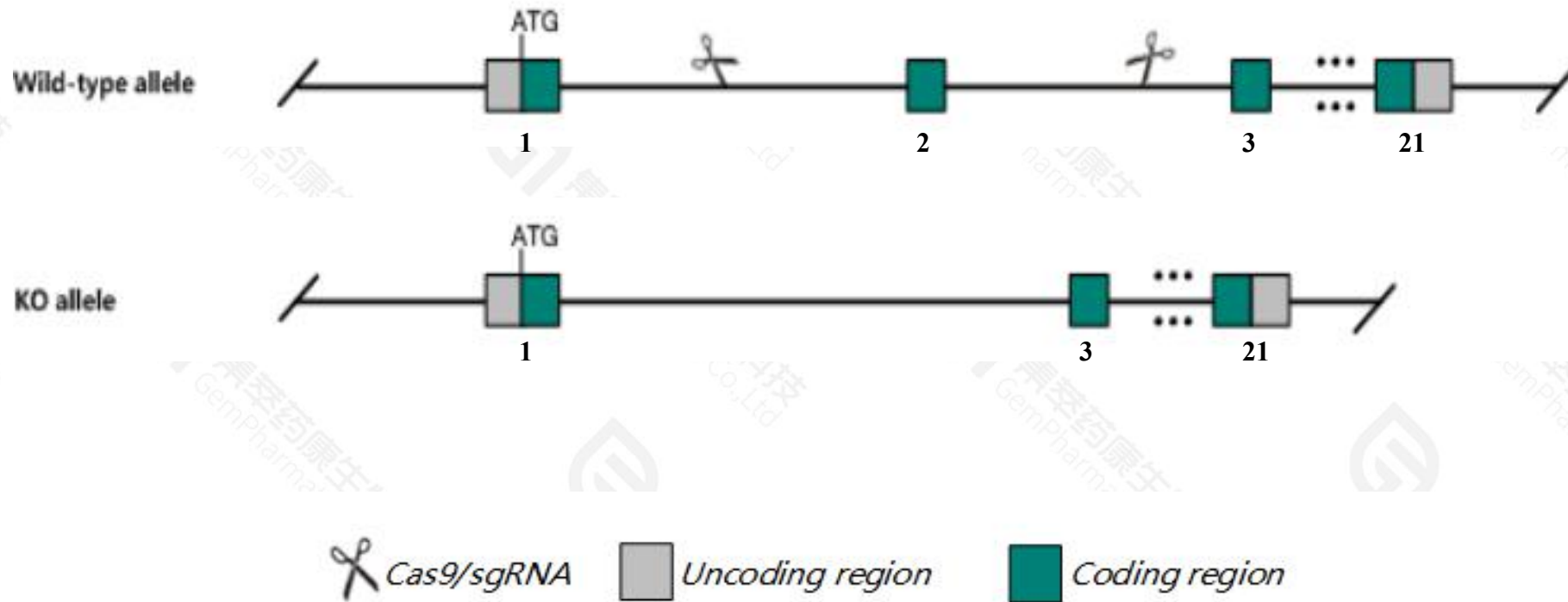
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Capn7* gene. The schematic diagram is as follows:



- The *Capn7* gene has 4 transcripts. According to the structure of *Capn7* gene, exon2 of *Capn7*-201(ENSMUST00000022451.14) transcript is recommended as the knockout region. The region contains 109bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Capn7* 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.

- According to the existing MGI data, mice homozygous for disruptions in this gene frequently die before weaning. Survivors display reduced body weight.
- The *Capn7* gene is located on the Chr14. 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.

Capn7 calpain 7 [Mus musculus (house mouse)]

Gene ID: 12339, updated on 17-Dec-2020

Summary



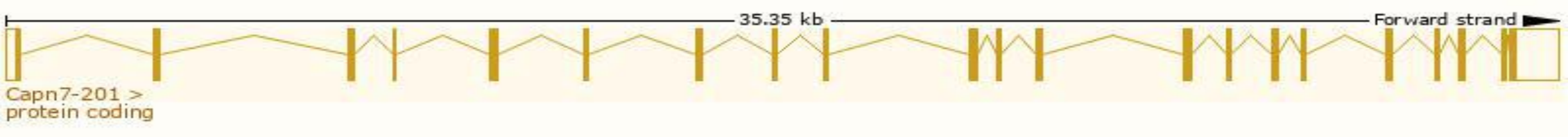
Official Symbol	Capn7 provided by MGI
Official Full Name	calpain 7 provided by MGI
Primary source	MGI:MGI:1338030
See related	Ensembl:ENSMUSG00000021893
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AU022319, Pa, PalBH
Expression	Ubiquitous expression in CNS E18 (RPKM 22.2), CNS E14 (RPKM 19.2) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

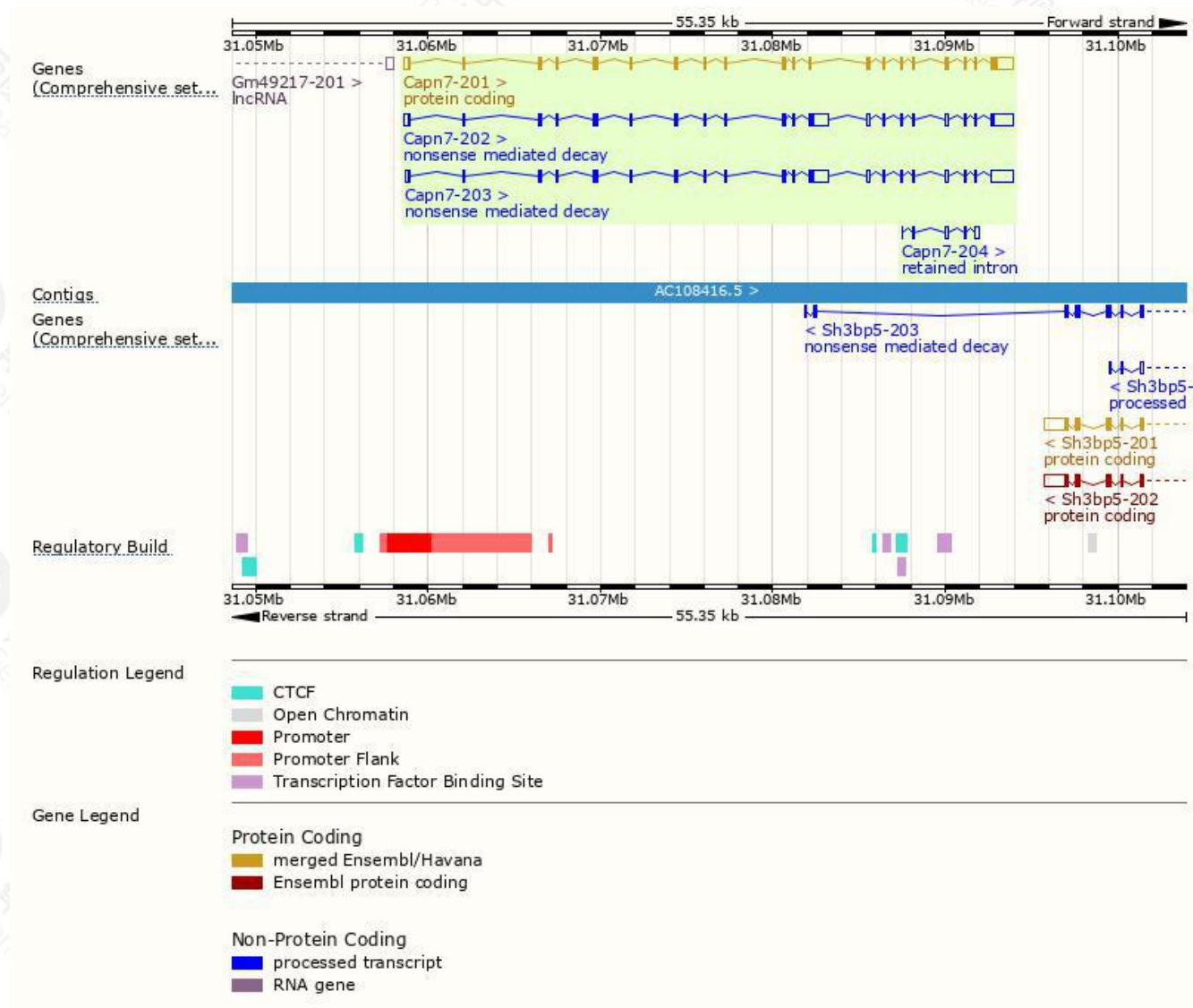
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Capn7-201	ENSMUST00000022451.14	3648	813aa	Protein coding	CCDS26911		TSL:1 , GENCODE basic , APPRIS P1 ,
Capn7-203	ENSMUST00000152182.2	4662	508aa	Nonsense mediated decay	-		TSL:1 ,
Capn7-202	ENSMUST00000143472.8	4613	508aa	Nonsense mediated decay	-		TSL:5 ,
Capn7-204	ENSMUST00000228237.2	614	No protein	Retained intron	-		

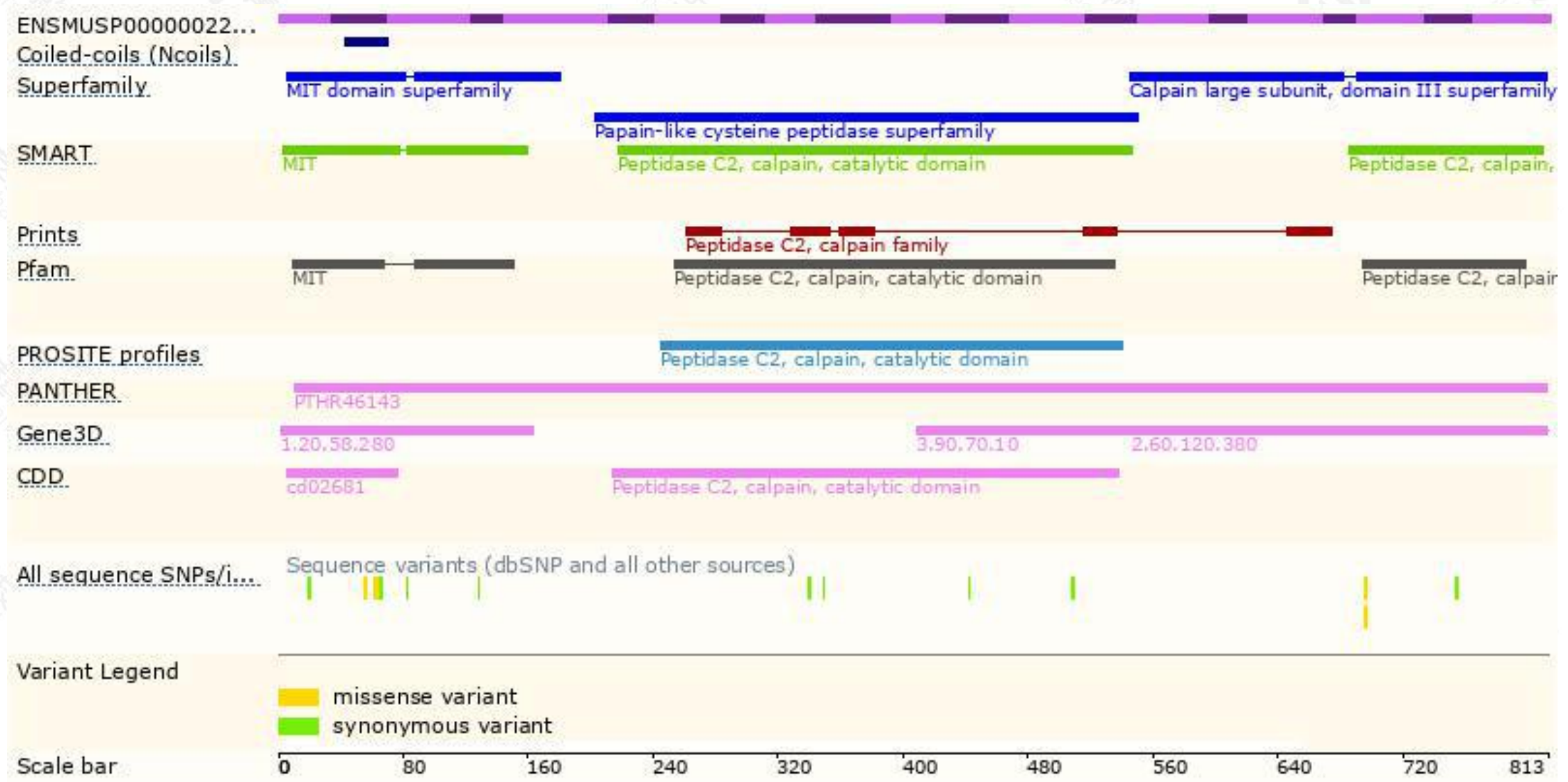
The strategy is based on the design of *Capn7-201* transcript,the transcription is shown below:



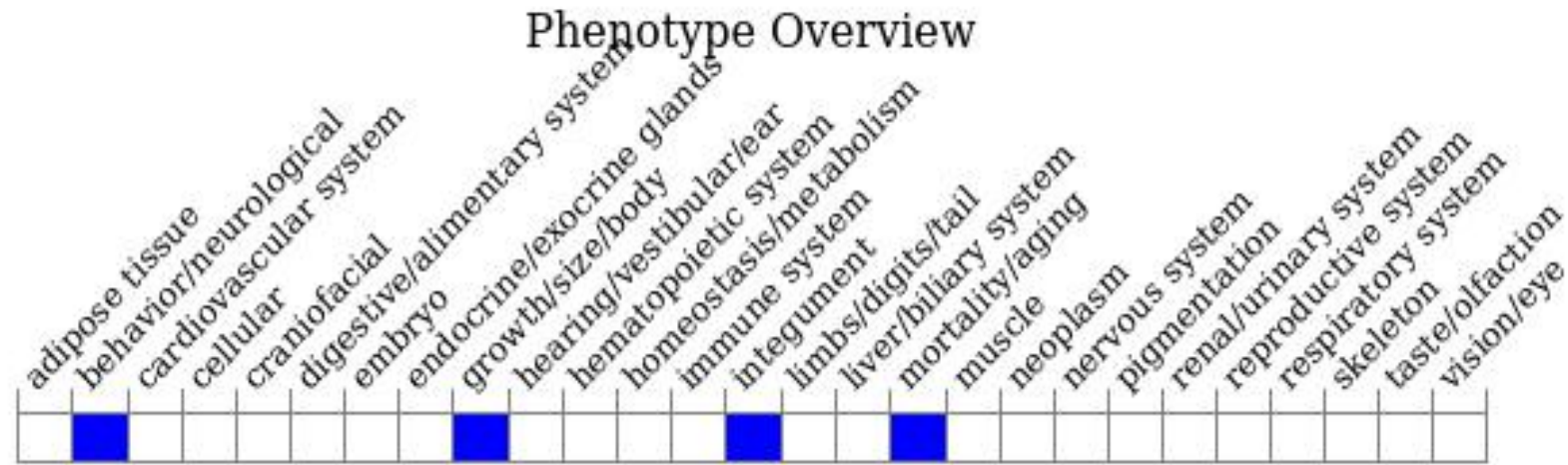
Genomic location distribution



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 disruptions in this gene frequently die before weaning.

Survivors display reduced body weight.

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

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