

Adap2 Cas9-KO Strategy

Designer: Xueting Zhang

Reviewer: Daohua Xu

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Project Overview

Project Name

Adap2

Project type

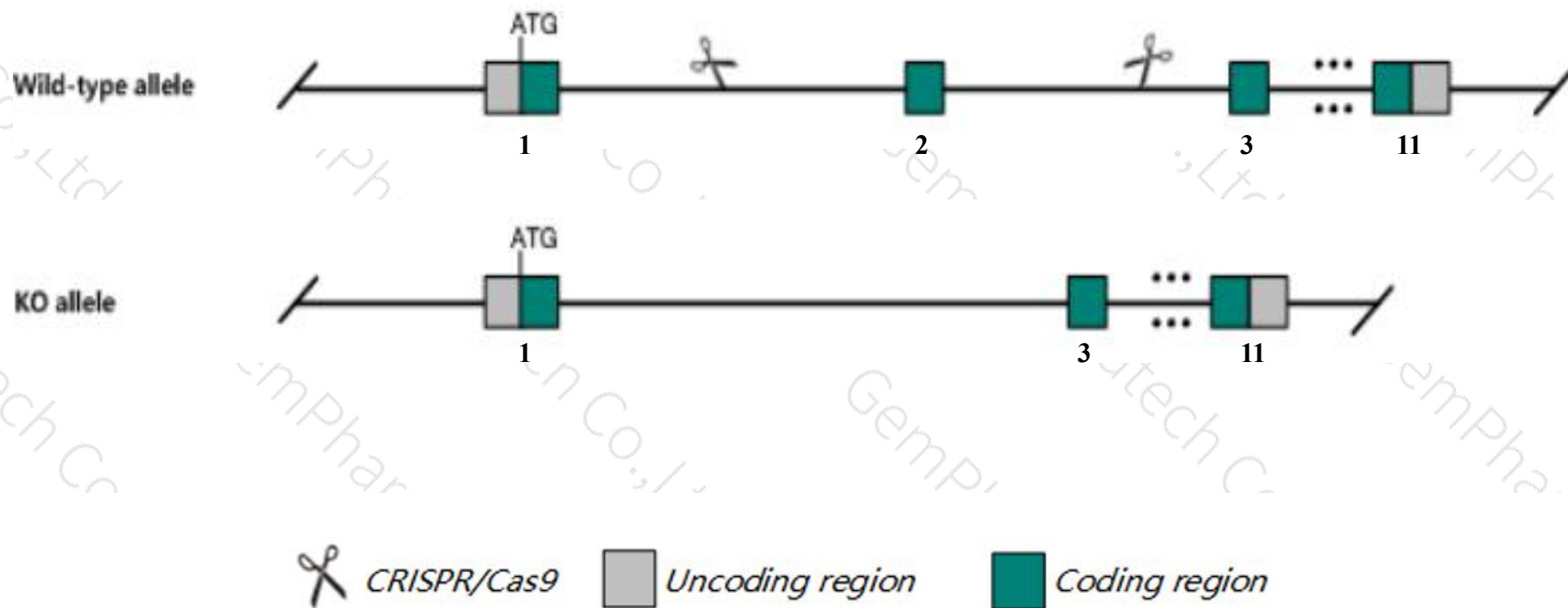
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Adap2* gene. The schematic diagram is as follows:



- The *Adap2* gene has 2 transcripts. According to the structure of *Adap2* gene, exon2 of *Adap2-201*(ENSMUST00000021050.13) transcript is recommended as the knockout region. The region contains 131bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Adap2* gene. The brief process is as follows: CRISPR/Cas9 system 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 *Adap2* gene is located on the Chr11. 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)

Adap2 ArfGAP with dual PH domains 2 [Mus musculus (house mouse)]

Gene ID: 216991, updated on 13-Mar-2020

Summary



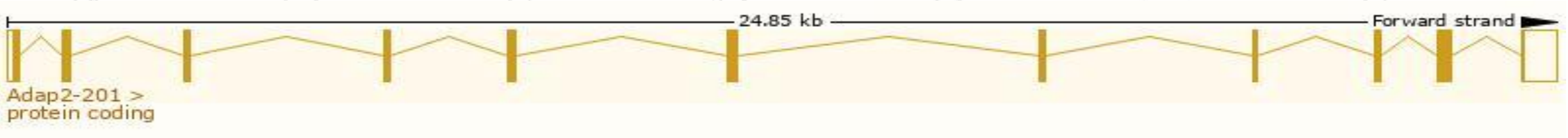
Official Symbol	Adap2 provided by MGI
Official Full Name	ArfGAP with dual PH domains 2 provided by MGI
Primary source	MGI:MGI:2663075
See related	Ensembl:ENSMUSG00000020709
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Centa2
Expression	Ubiquitous expression in kidney adult (RPKM 6.0), liver adult (RPKM 4.2) and 25 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

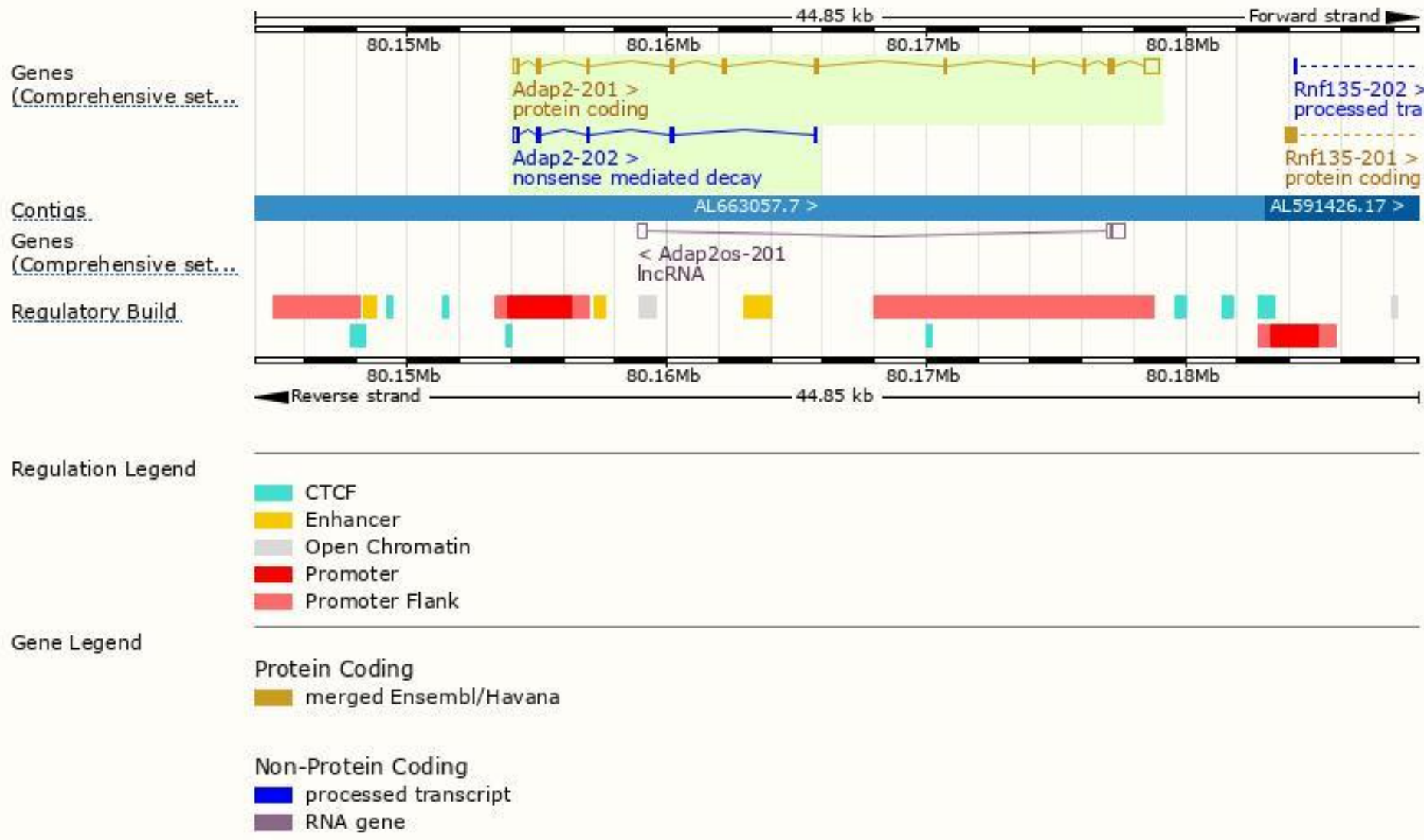
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adap2-201	ENSMUST00000021050.13	1785	381aa	Protein coding	CCDS25128	Q5SSK2 Q8R2V5	TSL:1 GENCODE basic APPRIS P1
Adap2-202	ENSMUST00000134118.1	601	133aa	Nonsense mediated decay	-	E9QAL8	TSL:2

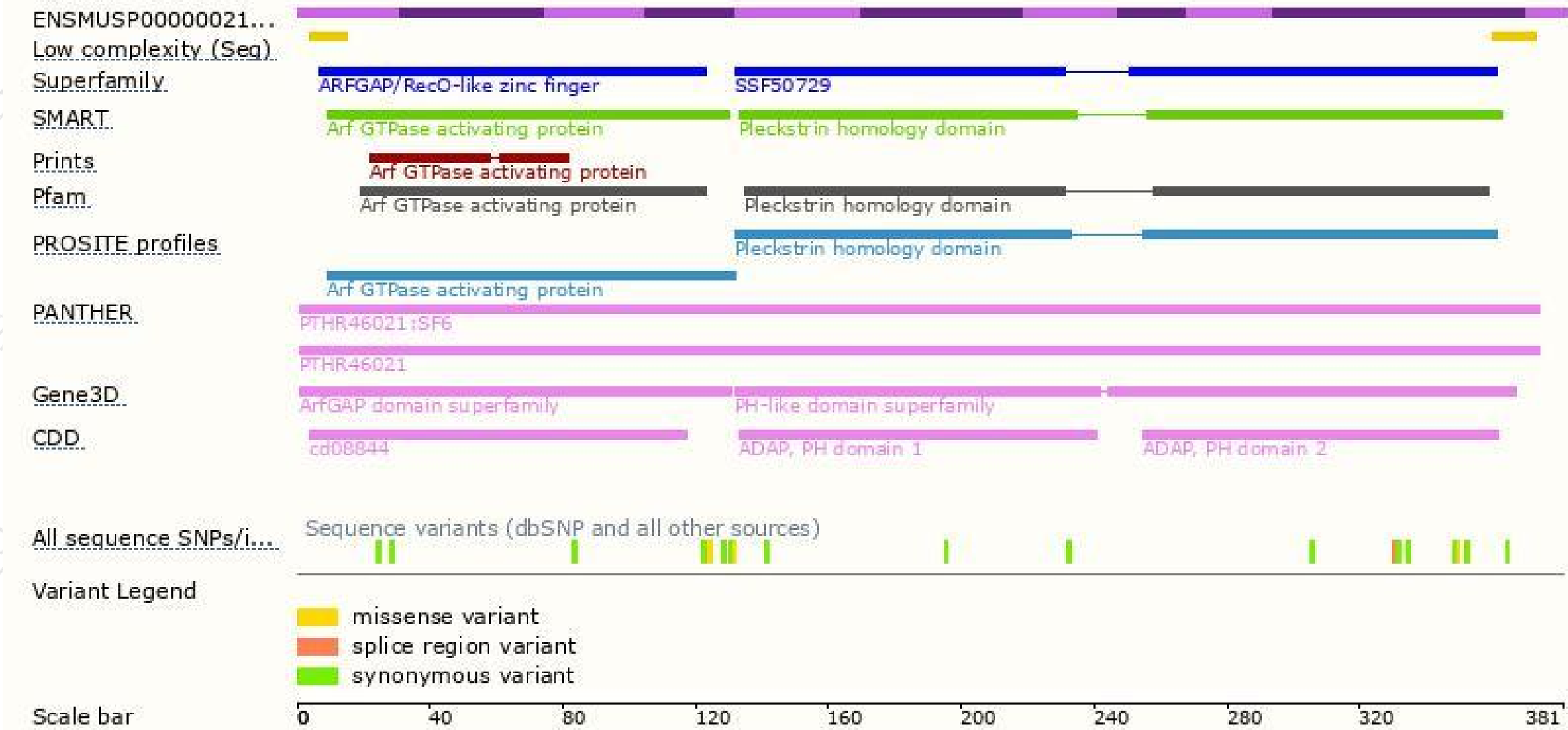
The strategy is based on the design of *Adap2-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

Tel: 400-9660890

