

Ajuba Cas9-KO Strategy

Designer: Shuang Zhang

Reviewer: Yun Li

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Project Overview

Project Name

Ajuba

Project type

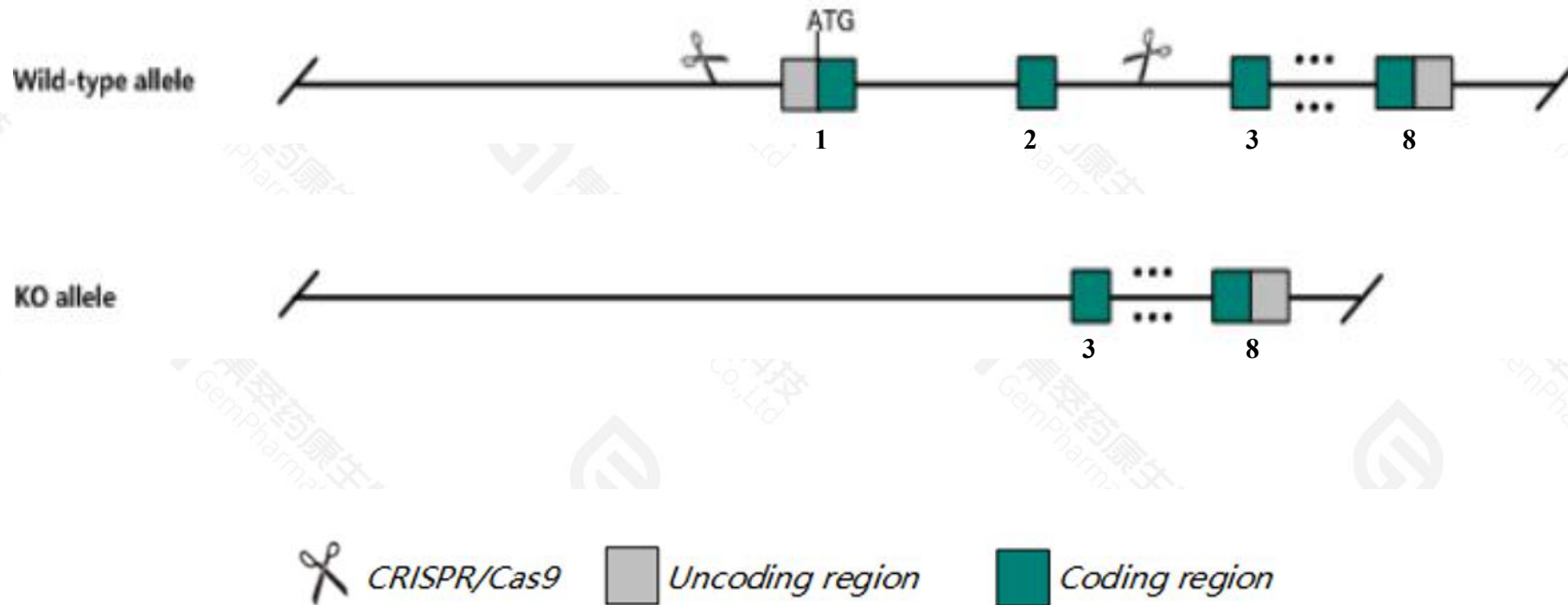
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Ajuba* gene has 2 transcripts. According to the structure of *Ajuba* gene, exon1-exon2 of *Ajuba*-201(ENSMUST00000054487.10) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ajuba* 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.

- According to the existing MGI data, homozygous null mice are viable and reach adulthood without any obvious phenotypes, however mouse embryonic fibroblasts exhibit impaired cell migration and abnormal lamellipodia production in vitro.
- The *Ajuba* 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.

Ajuba ajuba LIM protein [Mus musculus (house mouse)]

Gene ID: 16475, updated on 17-Nov-2020

Summary



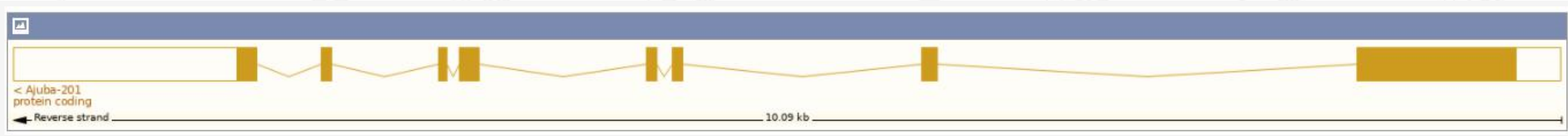
Official Symbol	Ajuba provided by MGI
Official Full Name	ajuba LIM protein provided by MGI
Primary source	MGI:MGI:1341886
See related	Ensembl:ENSMUSG00000022178
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Jub
Expression	Broad expression in adrenal adult (RPKM 26.2), ovary adult (RPKM 18.9) and 24 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

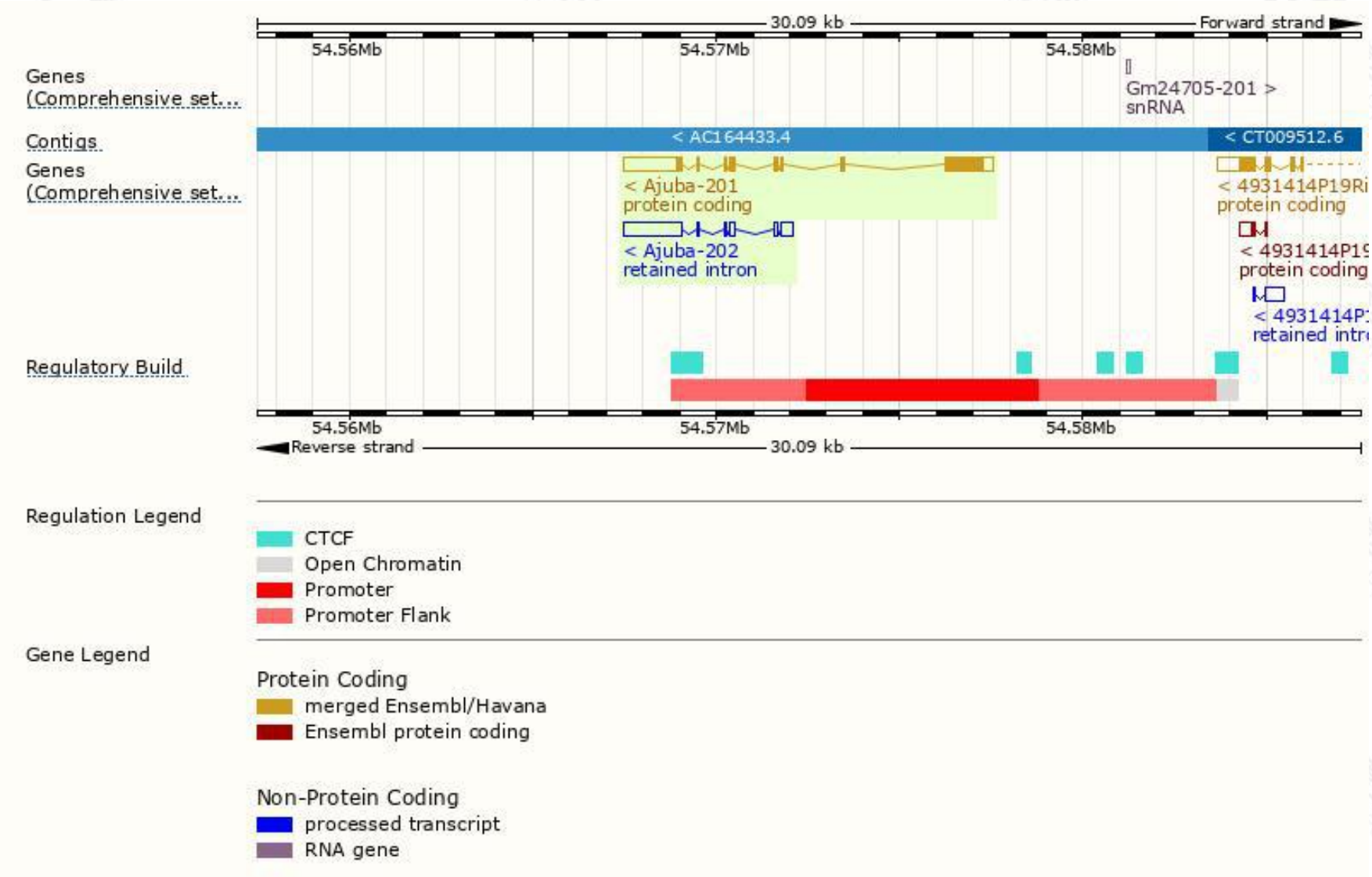
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ajuba-201	ENSMUST00000054487.10	3397	547aa	Protein coding	CCDS27093		TSL:1 , GENCODE basic , APPRIS P1 ,
Ajuba-202	ENSMUST00000226463.2	2214	No protein	Retained intron	-		

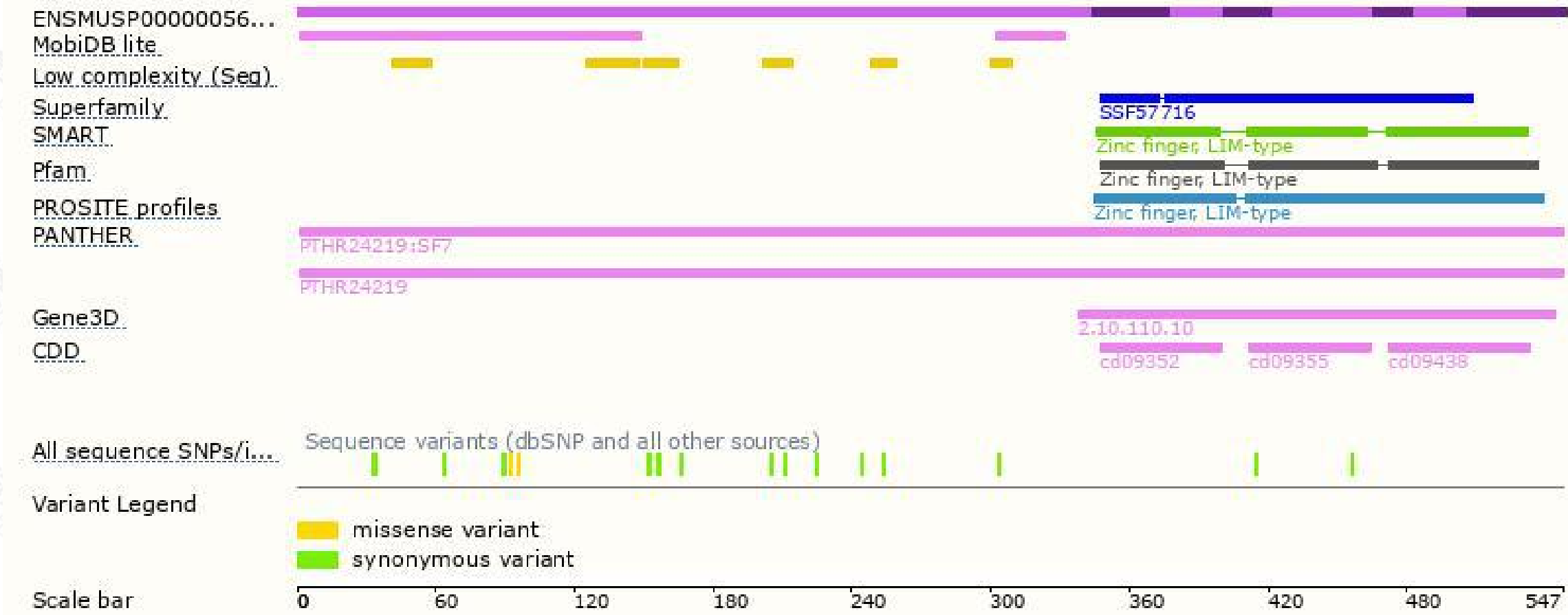
The strategy is based on the design of *Ajuba-201* transcript,the transcription is shown below:



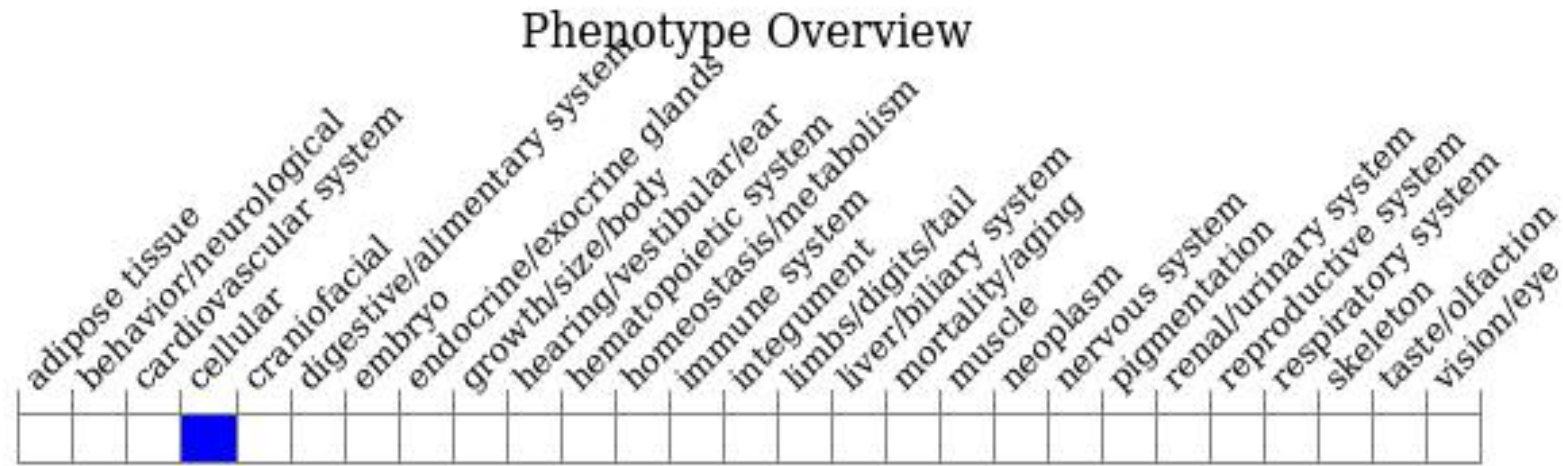
Genomic location distribution



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, homozygous null mice are viable and reach adulthood without any obvious phenotypes, however mouse embryonic fibroblasts exhibit impaired cell migration and abnormal lamellipodia production in vitro.

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

