

Arpc5l Cas9-KO Strategy

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Project Overview

Project Name

Arpc5l

Project type

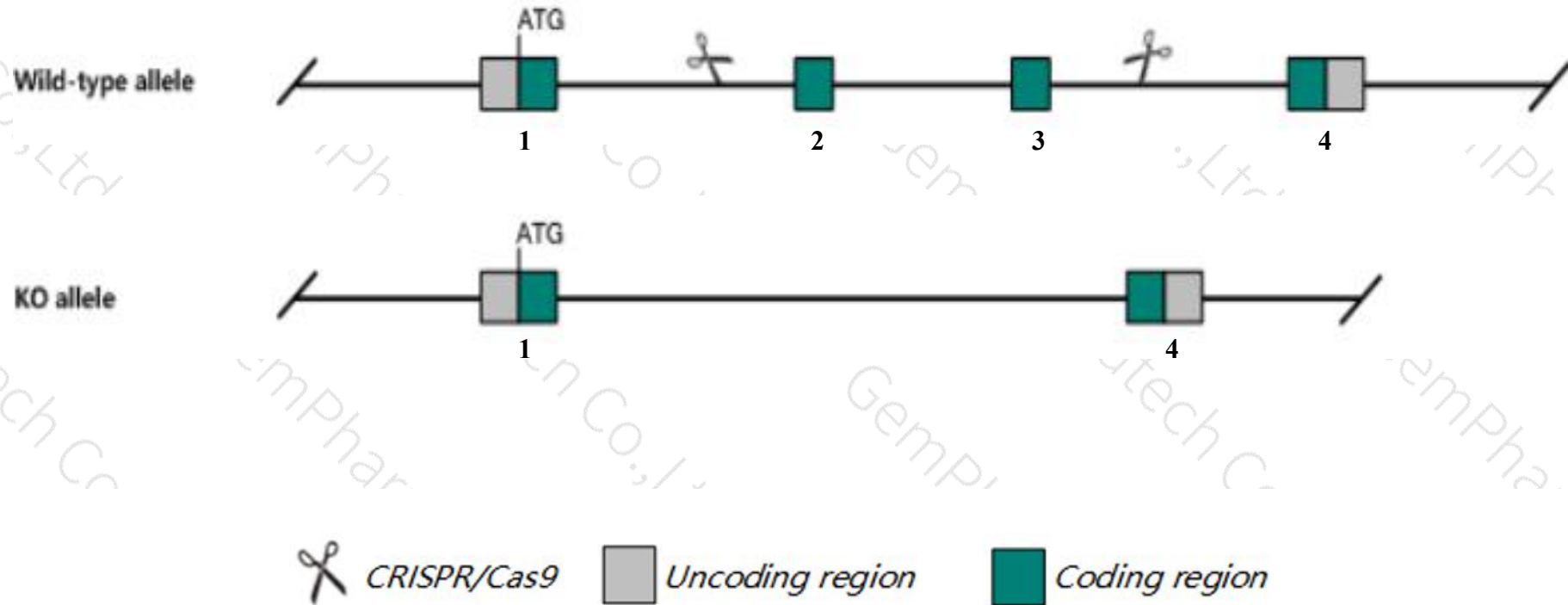
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Arpc5l* gene. The schematic diagram is as follows:



- The *Arpc5l* gene has 5 transcripts. According to the structure of *Arpc5l* gene, exon2-exon3 of *Arpc5l-202* (ENSMUST00000112862.6) transcript is recommended as the knockout region. The region contains 250bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Arpc5l* gene. The brief process is as follows: CRISPR/Cas9 system w

- The *Arpc5l* 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)

Arpc5l actin related protein 2/3 complex, subunit 5-like [*Mus musculus* (house mouse)]

Gene ID: 74192, updated on 3-May-2020

Summary

Official Symbol	Arpc5l provided by MGI
Official Full Name	actin related protein 2/3 complex, subunit 5-like provided by MGI
Primary source	MGI:MGI:1921442
See related	Ensembl:ENSMUSG00000026755
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	ARC16-2; AI852867; AW555592; AW742746; 2010015J01Rik
Expression	Ubiquitous expression in spleen adult (RPKM 20.9), testis adult (RPKM 20.3) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

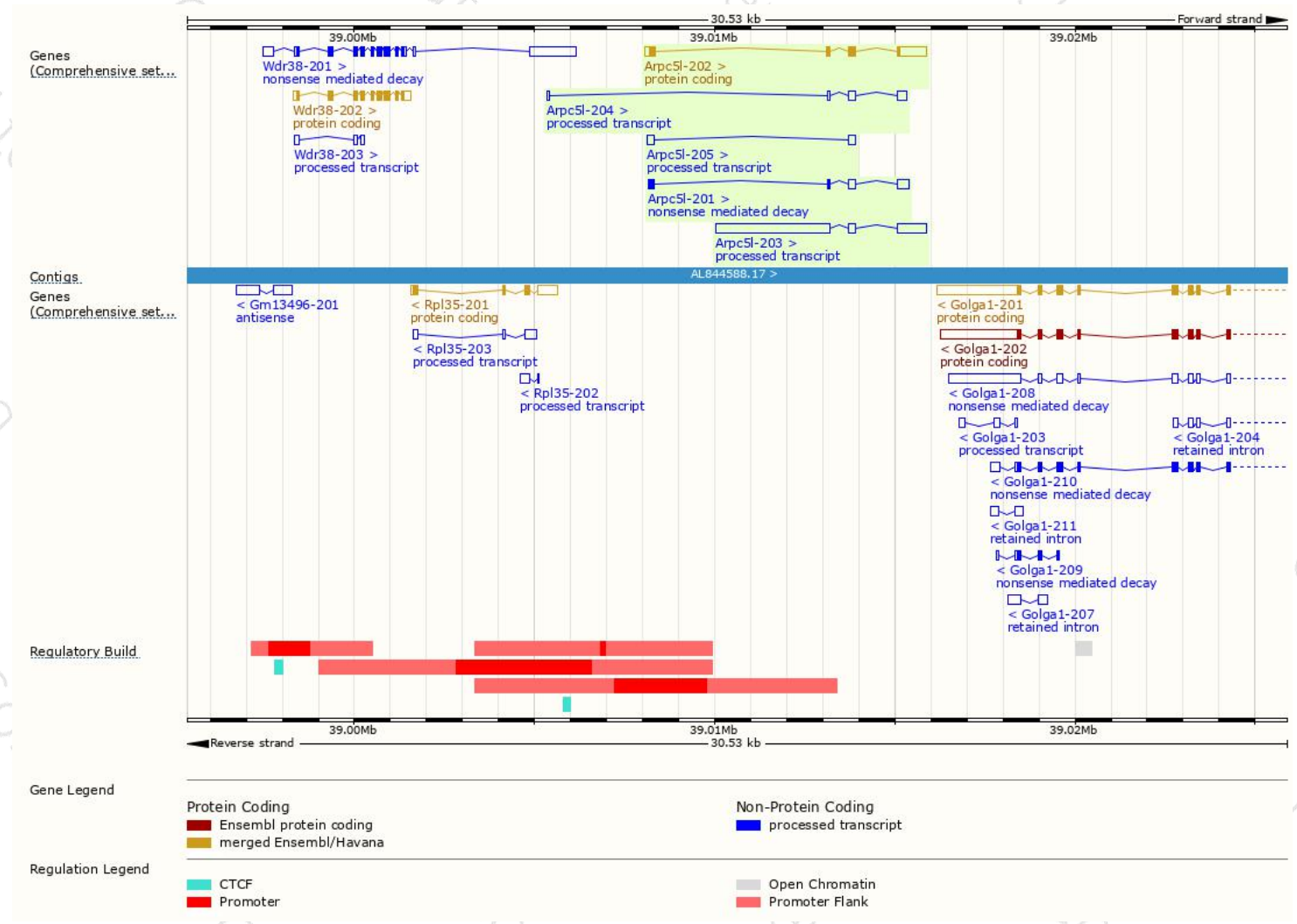
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Arpc5l-202	ENSMUST00000112862.6	1339	153aa	Protein coding	CCDS38120	Q9D898	TSL:1 GENCODE basic APPRIS P1
Arpc5l-201	ENSMUST00000090993.7	752	65aa	Nonsense mediated decay	-	A0A0R3P9C9	TSL:2
Arpc5l-203	ENSMUST00000135049.1	4167	No protein	Processed transcript	-	-	TSL:1
Arpc5l-204	ENSMUST00000141467.7	579	No protein	Processed transcript	-	-	TSL:3
Arpc5l-205	ENSMUST00000204825.2	361	No protein	Processed transcript	-	-	TSL:2

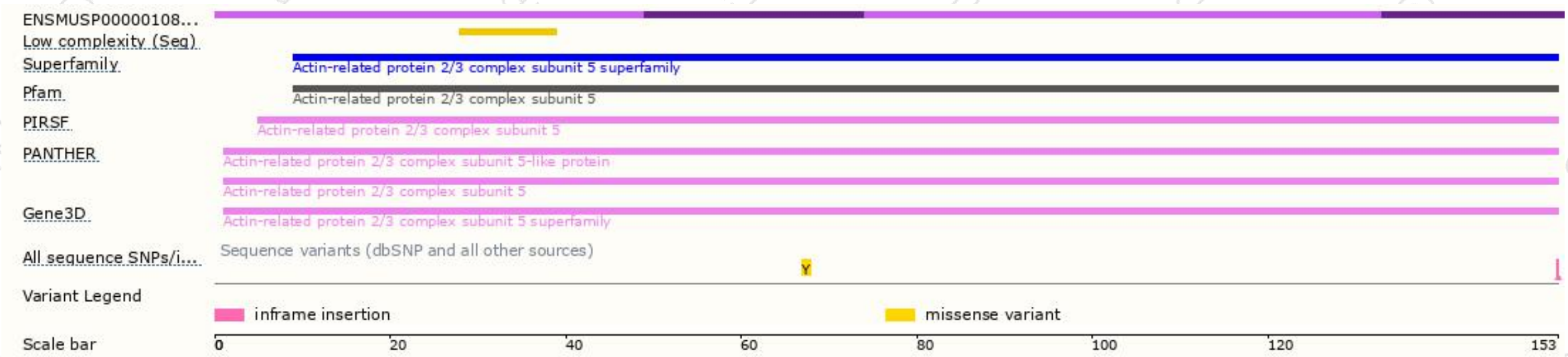
The strategy is based on the design of *Arpc5l-202* transcript, the transcription is shown below



Genomic location distribution



Protein domain



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

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