

Farp1 Cas9-KO Strategy

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Design Date: 2020-8-10

Project Overview

Project Name

Farp1

Project type

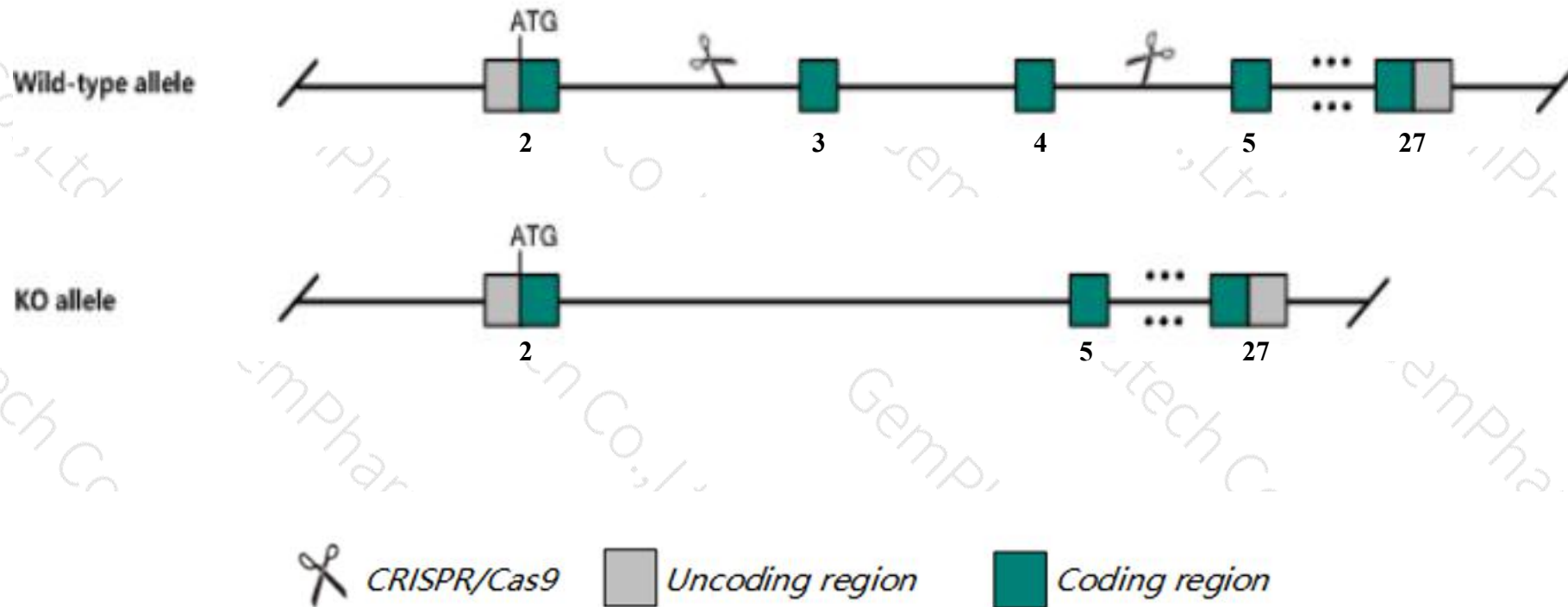
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Farp1* gene. The schematic diagram is as follows:



- The *Farp1* gene has 4 transcripts. According to the structure of *Farp1* gene, exon3-exon4 of *Farp1*-201(ENSMUST00000026635.7) transcript is recommended as the knockout region. The region contains 148bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Farp1* 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 *Farp1* 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.
- Transcript *Farp1*-203 may not be affected.
- 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)

Farp1 FERM, RhoGEF (Arhgef) and pleckstrin domain protein 1 (chondrocyte-derived) [Mus musculus (house mouse)]

Gene ID: 223254, updated on 13-Mar-2020

Summary



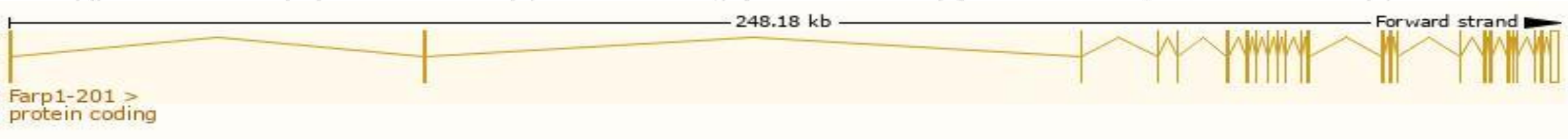
Official Symbol	Farp1 provided by MGI
Official Full Name	FERM, RhoGEF (Arhgef) and pleckstrin domain protein 1 (chondrocyte-derived) provided by MGI
Primary source	MGI:MGI:2446173
See related	Ensembl:ENSMUSG00000025555
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AW228844, BC030329, Cdep
Expression	Ubiquitous expression in limb E14.5 (RPKM 25.3), CNS E18 (RPKM 22.0) and 27 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

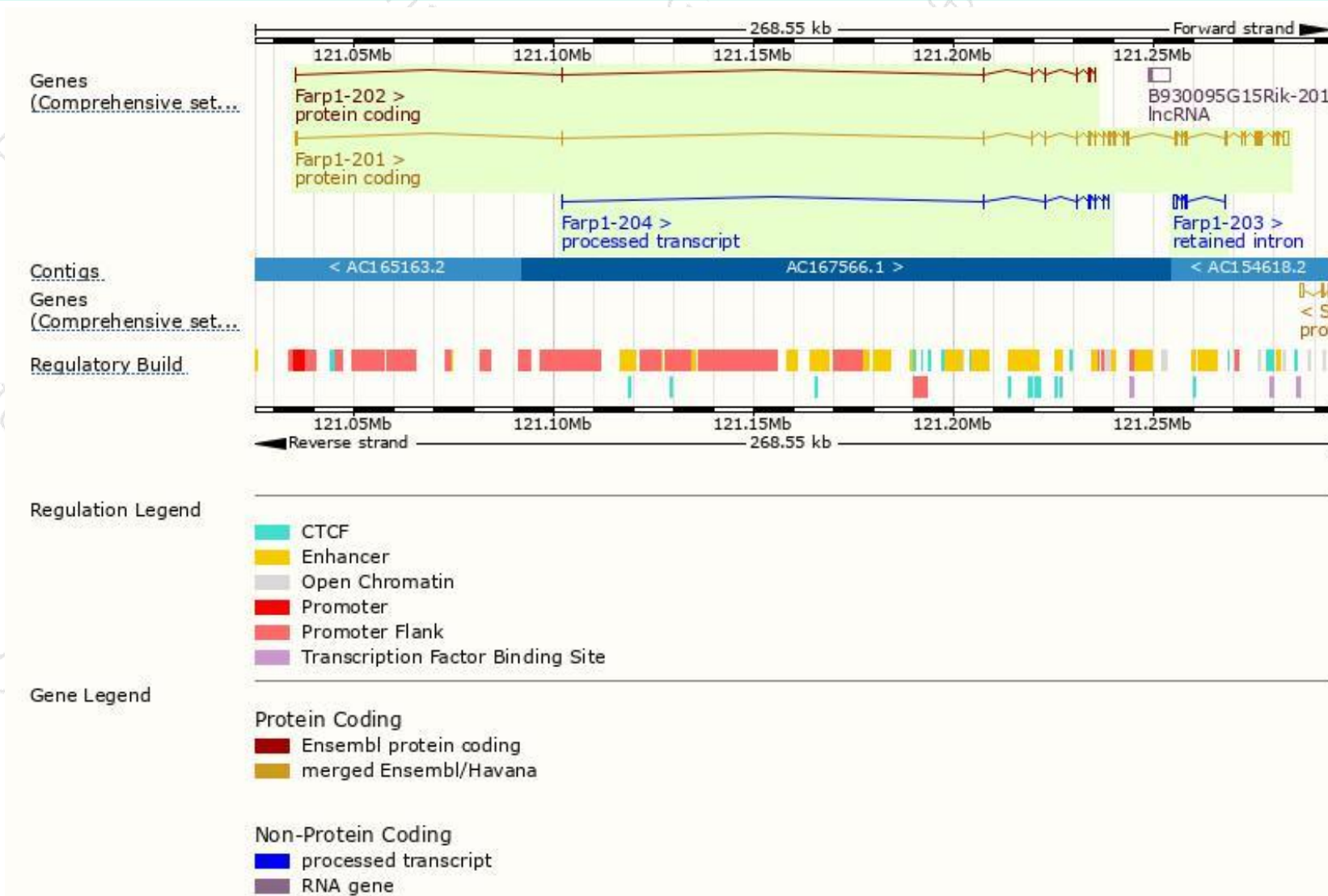
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Farp1-201	ENSMUST00000026635.7	4885	1048aa	Protein coding	CCDS37015	F8VPU2	TSL:5 GENCODE basic APPRIS P1
Farp1-202	ENSMUST00000135010.7	797	211aa	Protein coding	-	E9Q805	CDS 3' incomplete TSL:5
Farp1-204	ENSMUST00000153607.1	777	No protein	Processed transcript	-	-	TSL:3
Farp1-203	ENSMUST00000137971.1	961	No protein	Retained intron	-	-	TSL:3

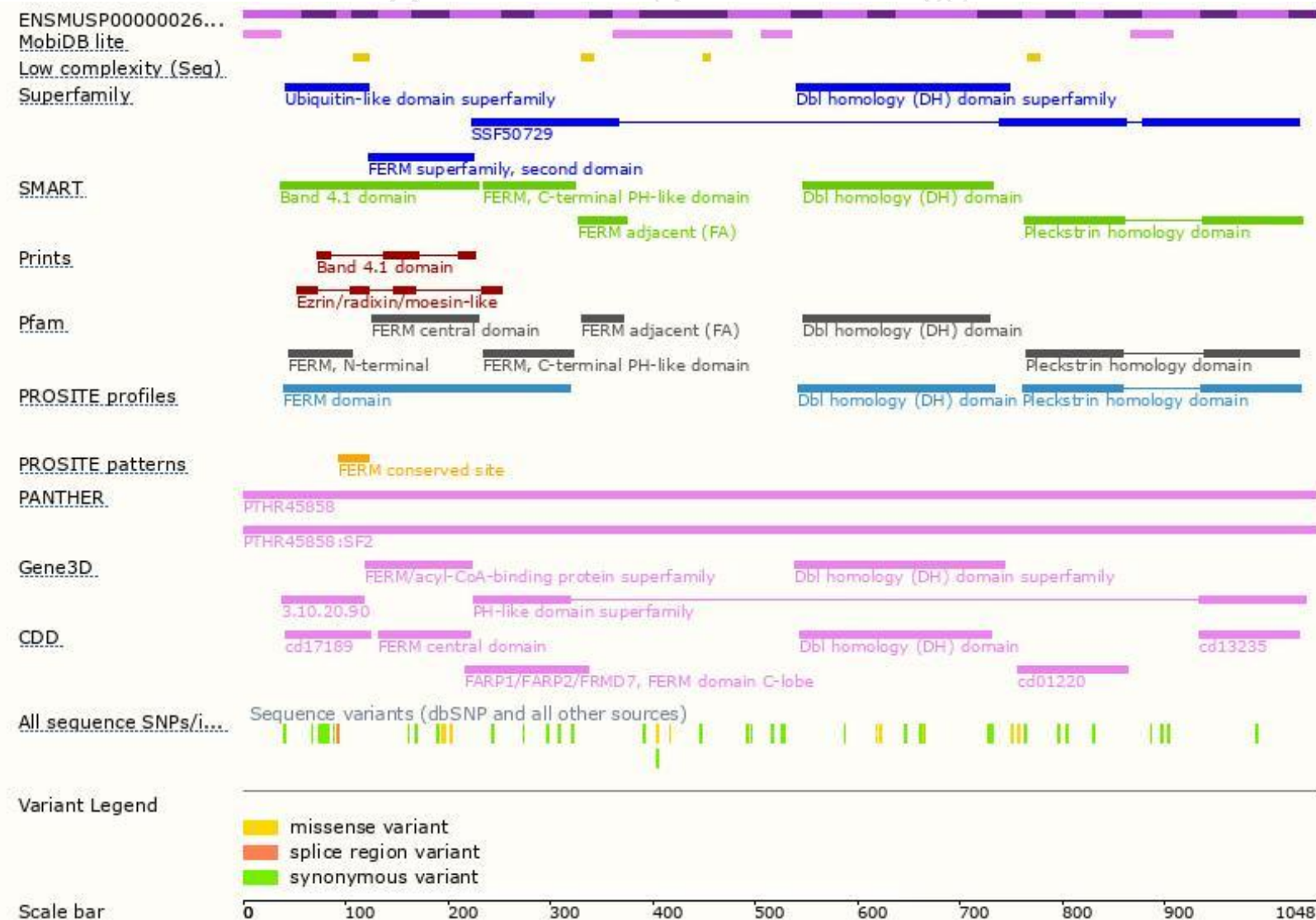
The strategy is based on the design of *Farp1-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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