

Rragd Cas9-CKO Strategy

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Project Overview

Project Name

Rragd

Project type

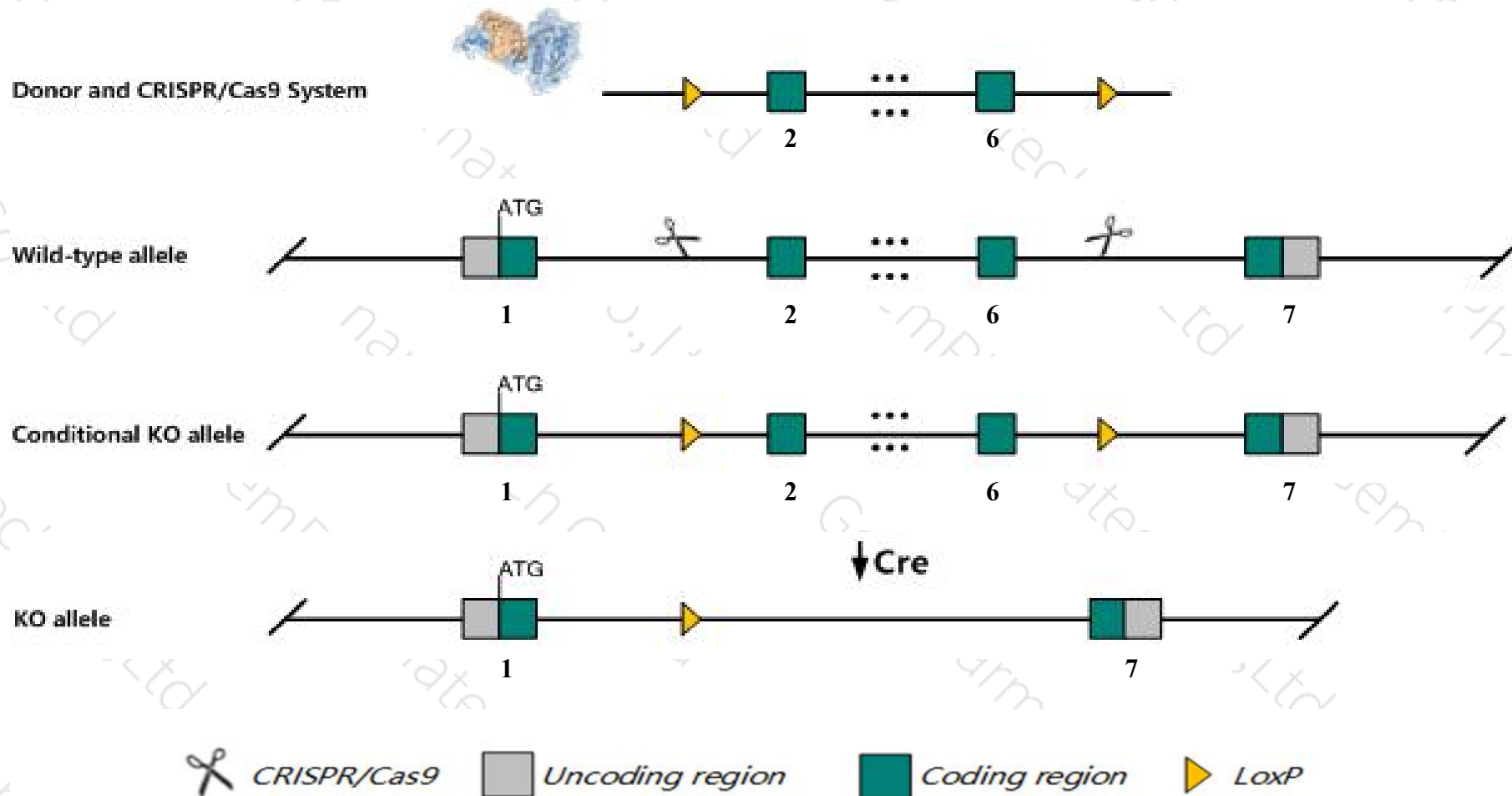
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



- The *Rragd* gene has 4 transcripts. According to the structure of *Rragd* gene, exon2-exon6 of *Rragd-203* (ENSMUST00000098190.9) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rragd* gene. The brief process is as follows: CRISPR/Cas9 system and Donor 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 flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- *Gm11943-201* gene may be destroyed.
- The *Rragd* gene is located on the Chr4. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Rragd Ras-related GTP binding D [*Mus musculus* (house mouse)]

Gene ID: 52187, updated on 8-Dec-2019

Summary

Official Symbol	Rragd provided by MGI
Official Full Name	Ras-related GTP binding D provided by MGI
Primary source	MGI:MGI:1098604
See related	Ensembl:ENSMUSG00000028278
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	AI467523; D4ErtD174e; 5730543C08Rik; C030003H22Rik
Expression	Broad expression in frontal lobe adult (RPKM 27.7), cortex adult (RPKM 13.1) and 17 other tissues See more
Orthologs	human all

Genomic context

Location: 4 A5; 4 14.57 cM

Exon count: 8

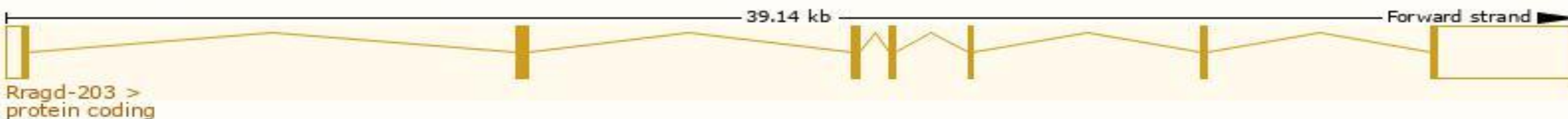
See Rragd in [Genome Data Viewer](#)

Transcript information (Ensembl)

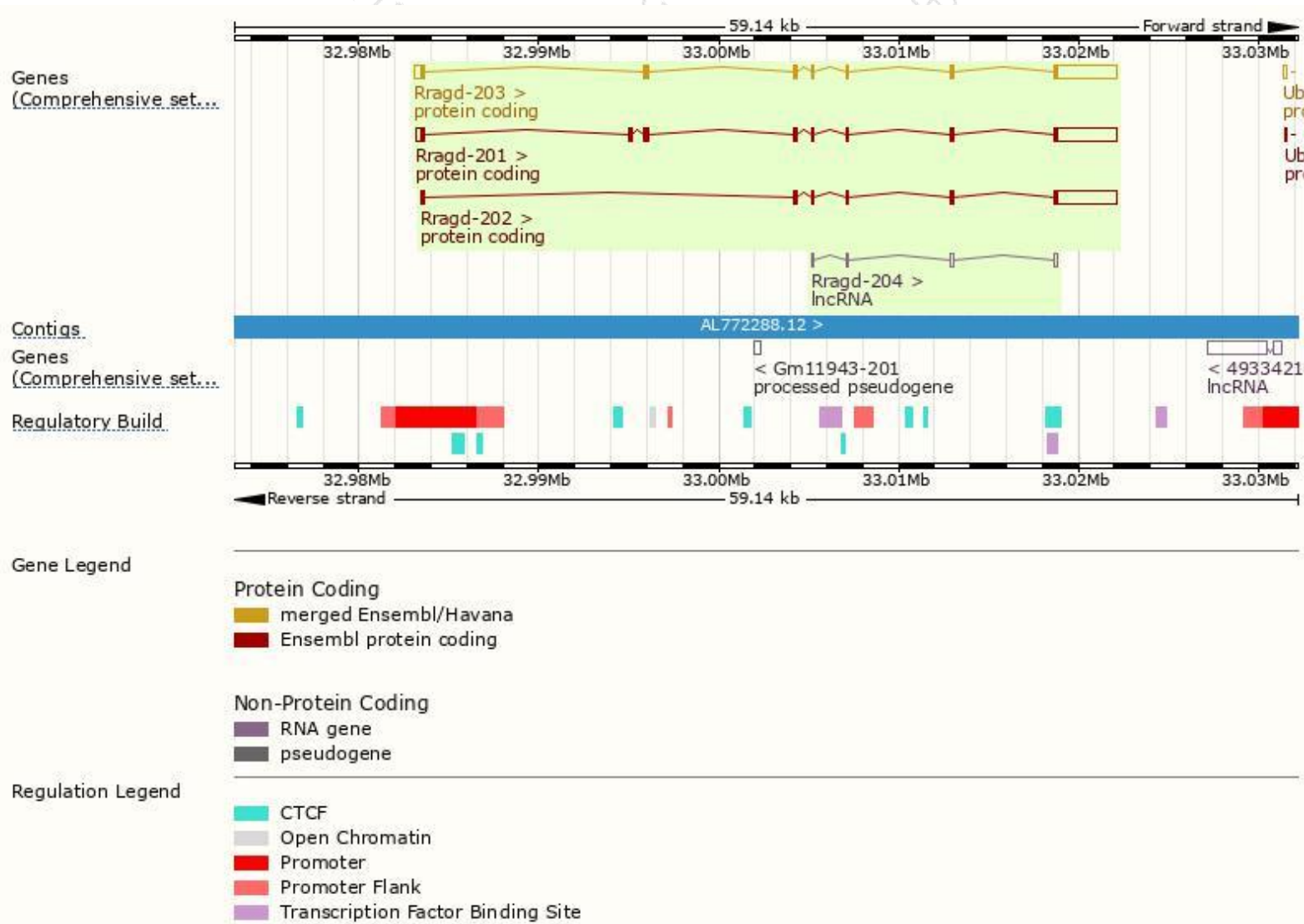
The gene has 4 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rragd-203	ENSMUST00000098190.9	5008	399aa	Protein coding	CCDS38704	B1AWT2	TSL:2 GENCODE basic APPRIS P1
Rragd-201	ENSMUST00000029946.13	5055	454aa	Protein coding	-	B1AWT3	TSL:5 GENCODE basic
Rragd-202	ENSMUST00000084747.5	4295	279aa	Protein coding	-	B1AWT4	TSL:5 GENCODE basic
Rragd-204	ENSMUST00000136792.1	452	No protein	lncRNA	-	-	TSL:3

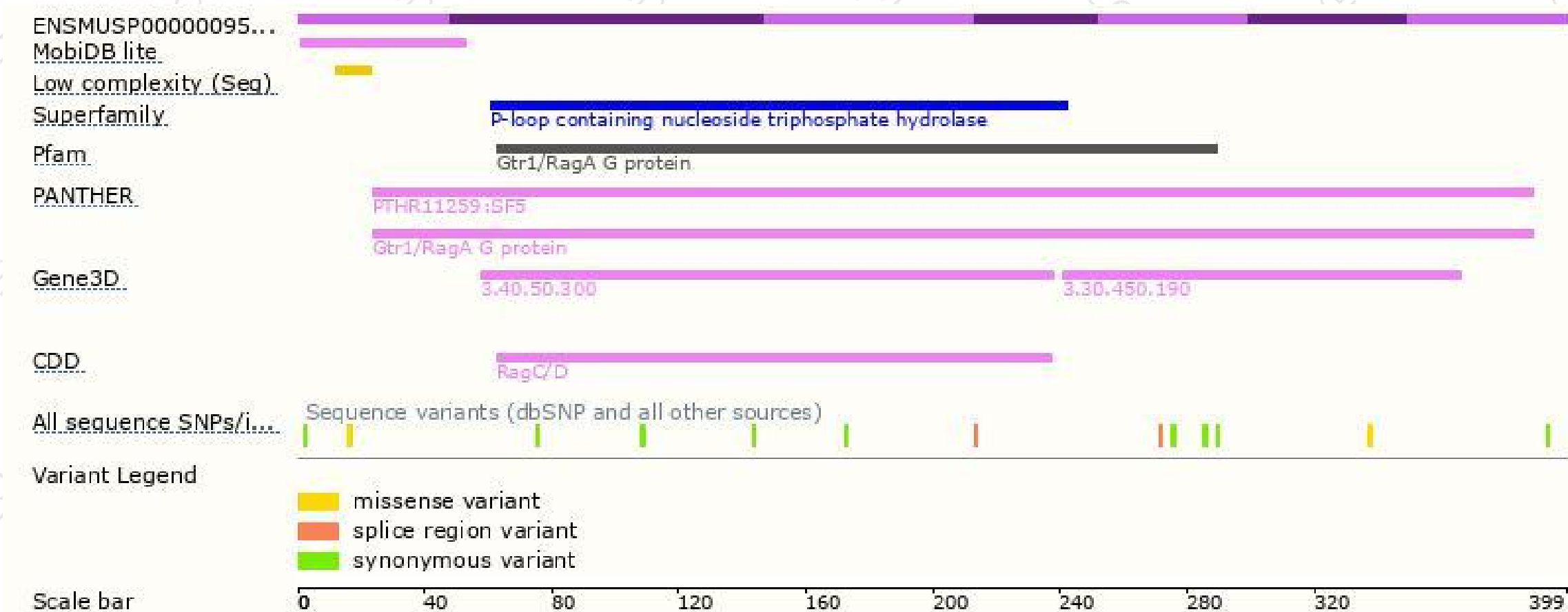
The strategy is based on the design of *Rragd-203* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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