

***Rab2a* Cas9-CKO Strategy**

Designer:

Ruirui Zhang

Reviewer:

Huimin Su

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Project Overview

Project Name

Rab2a

Project type

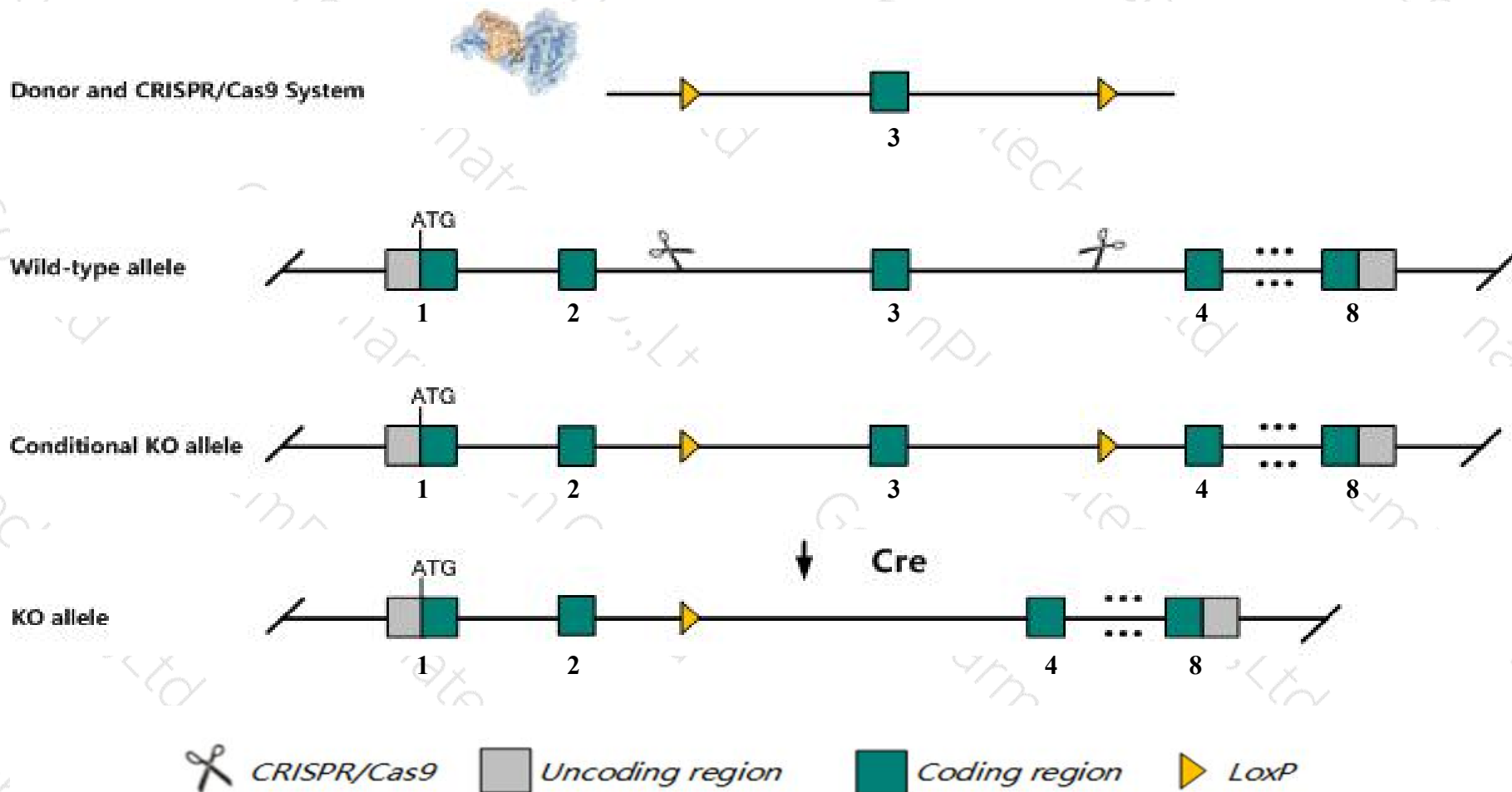
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rab2a* gene. The schematic diagram is as follows:



- The *Rab2a* gene has 1 transcript. According to the structure of *Rab2a* gene, exon3 of *Rab2a-201* (ENSMUST00000060232.7) transcript is recommended as the knockout region. The region contains 68bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rab2a* 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.

- According to the existing MGI data, mice homozygous for a knock-out allele exhibit preweaning lethality.
- The *Rab2a* 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)

Rab2a RAB2A, member RAS oncogene family [*Mus musculus* (house mouse)]

Gene ID: 59021, updated on 3-Nov-2019

Summary

Official Symbol Rab2a provided by [MGI](#)

Official Full Name RAB2A, member RAS oncogene family provided by [MGI](#)

Primary source [MGI:MGI:1928750](#)

See related [Ensembl:ENSMUSG000000047187](#)

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 Rab2; C80220; 9330148M11Rik

Expression Ubiquitous expression in CNS E18 (RPKM 79.3), adrenal adult (RPKM 75.7) and 28 other tissues [See more](#)

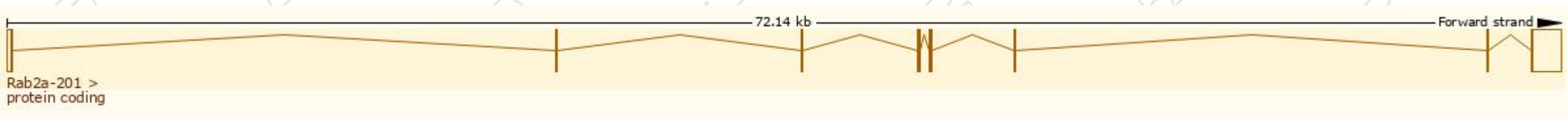
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

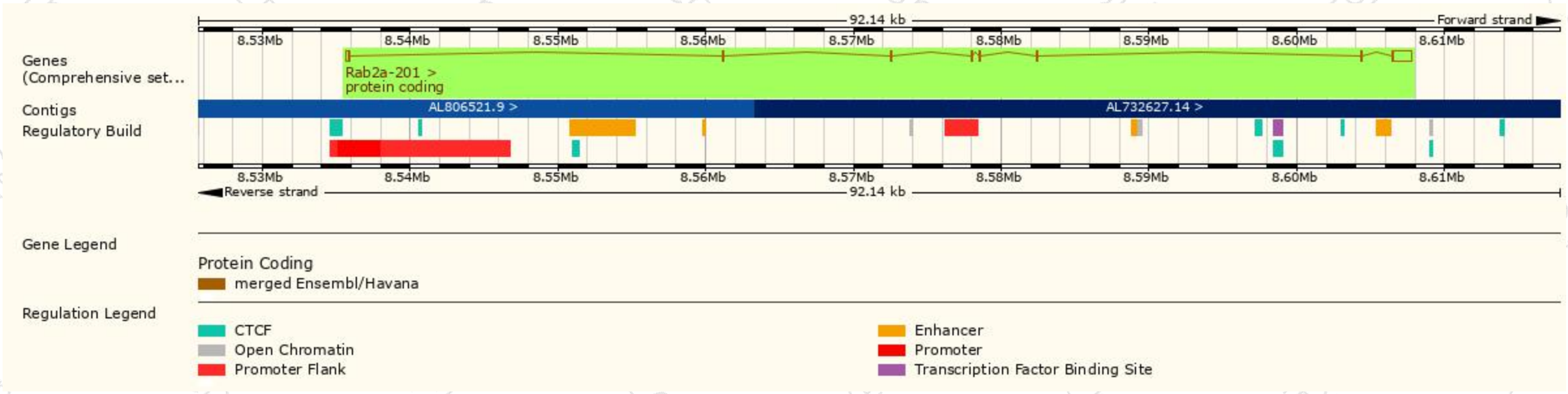
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rab2a-201	ENSMUST00000060232.7	2133	212aa	Protein coding	CCDS17955	P53994 Q0PD65	TSL:1 Gencode basic APPRIS P1

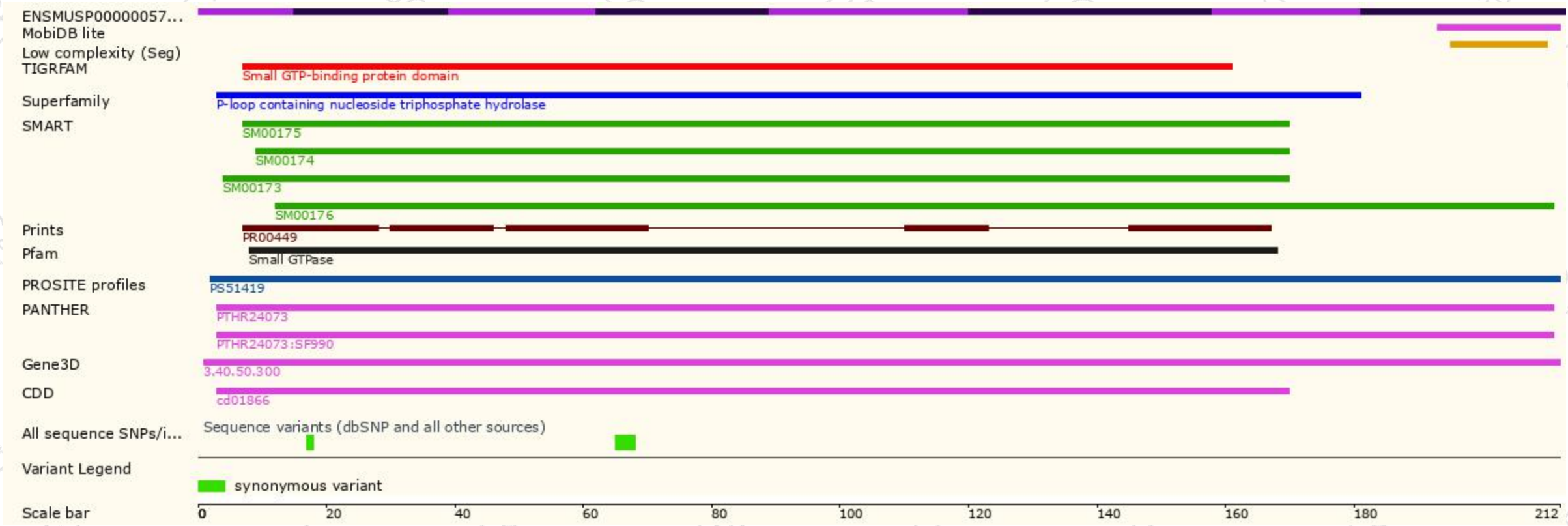
The strategy is based on the design of *Rab2a-201* transcript, the transcription is shown below:



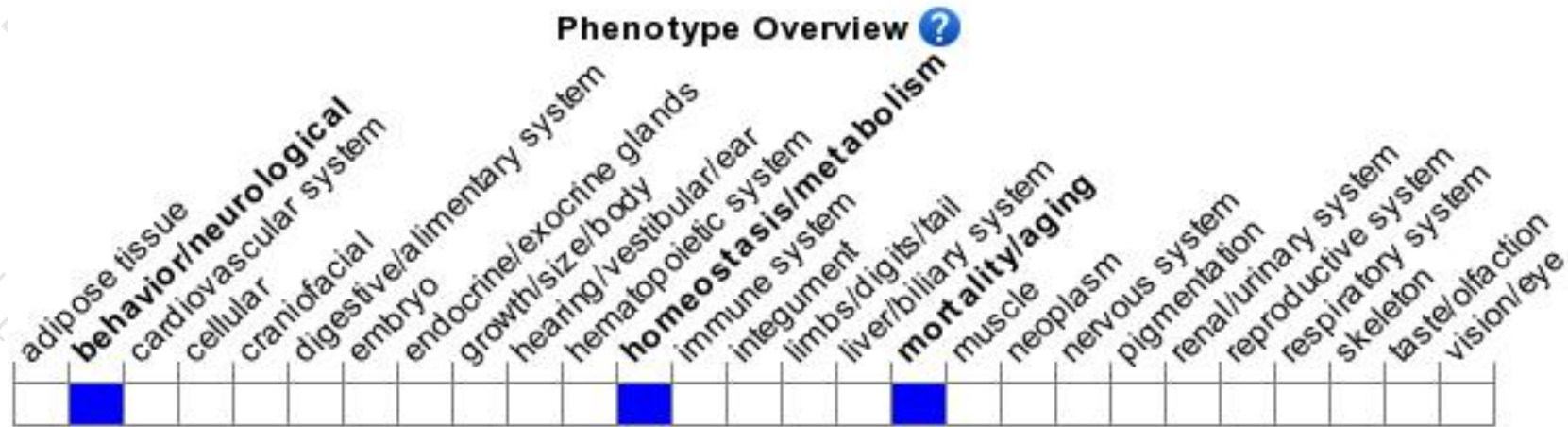
Genomic location distribution



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, mice homozygous for a knock-out allele exhibit preweaning lethality.

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

Tel: 400-9660890

