

***Rab2a* Cas9-KO Strategy**

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Reviewer:

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Project Overview

Project Name

Rab2a

Project type

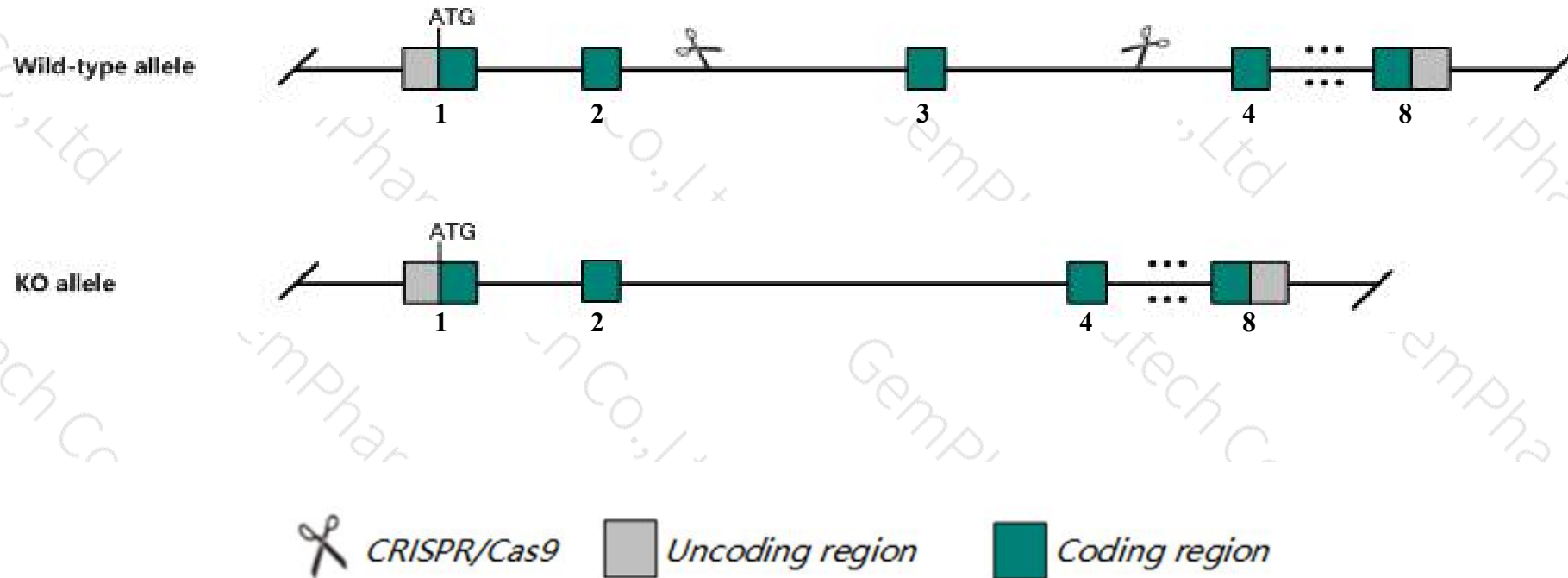
Cas9-KO

Strain background

C57BL/6JGpt

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

- 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

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

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

Summary

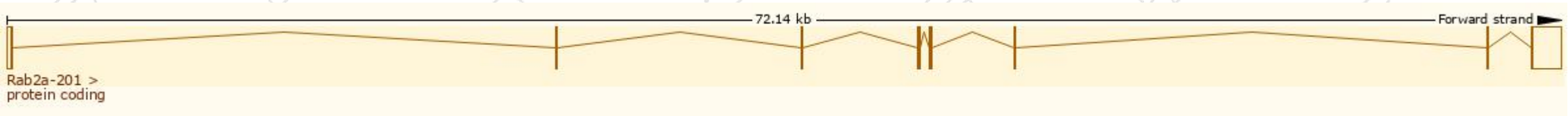
Official Symbol	Rab2a provided by MGI
Official Full Name	RAB2A, member RAS oncogene family provided by MGI
Primary source	MGI:MGI:1928750
See related	Ensembl:ENSMUSG000000047187
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; 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)

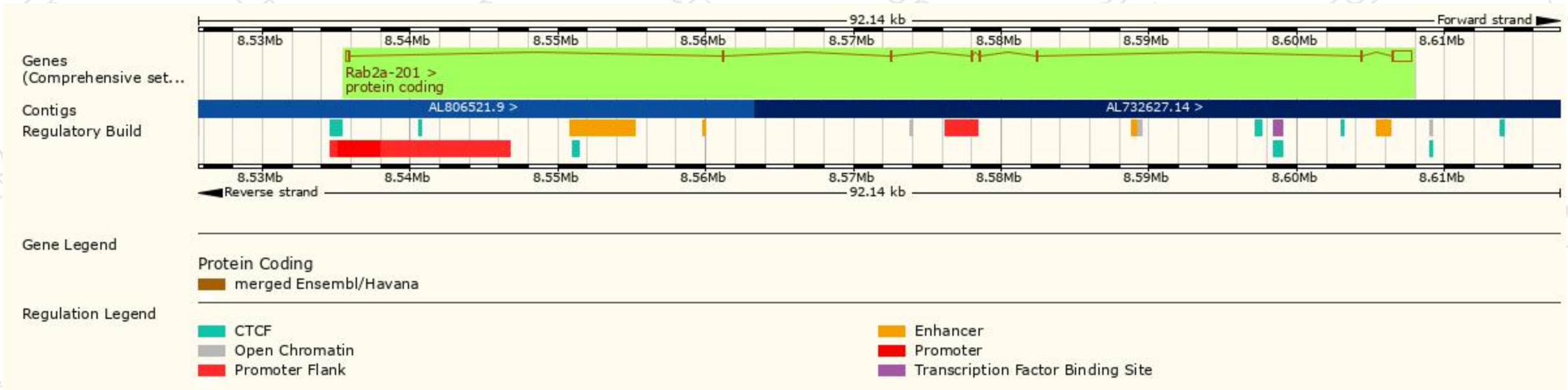
The gene has 1 transcript, and the transcript is shown below:

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

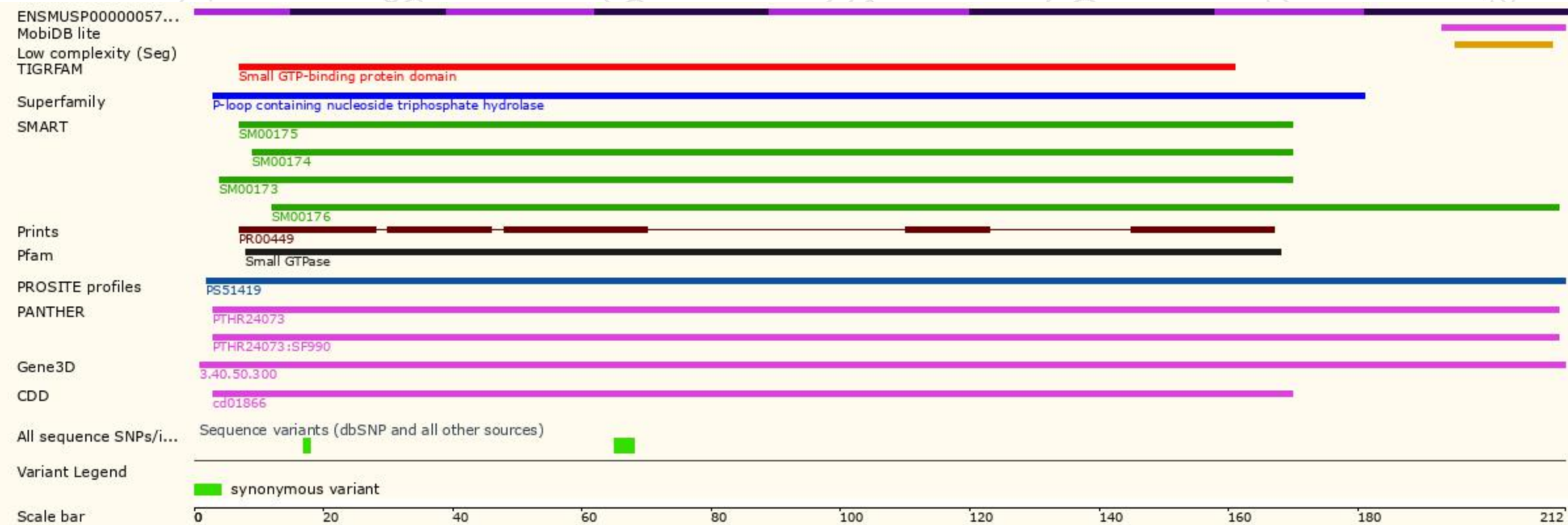
The strategy is based on the design of *Rab2a-201* transcript, the transcription is shown below:



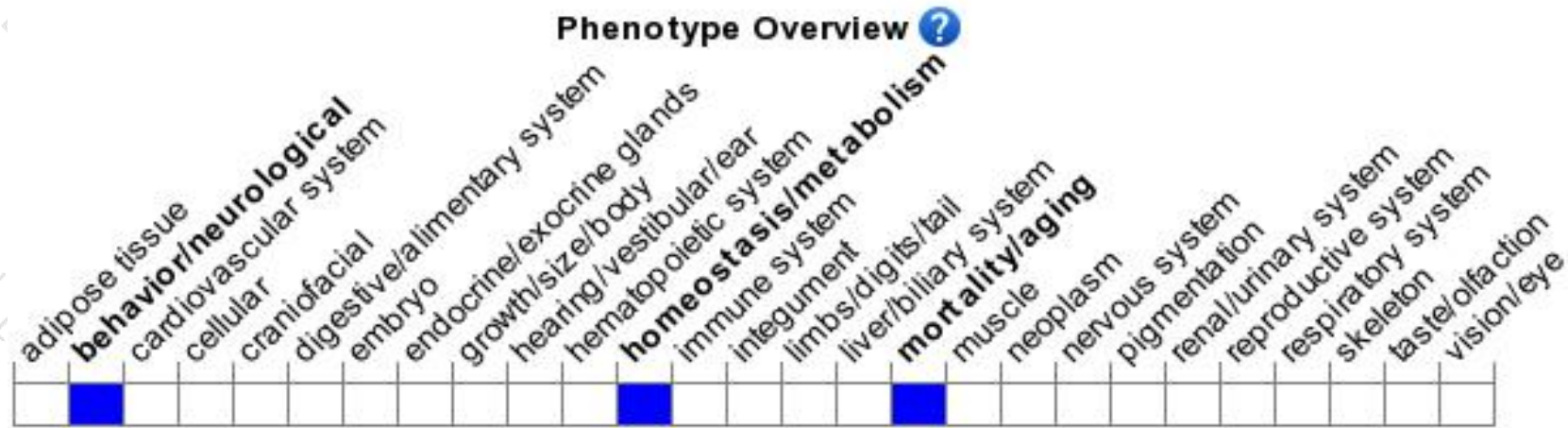
Genomic location distribution



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.

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