

***Rab5a* Cas9-KO Strategy**

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Reviewer:

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Project Overview

Project Name

Rab5a

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rab5a* gene. The schematic diagram is as follows:



- The *Rab5a* gene has 2 transcripts. According to the structure of *Rab5a* gene, exon3-exon4 of *Rab5a-201* (ENSMUST00000017975.6) transcript is recommended as the knockout region. The region contains 275bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rab5a* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Rab5a* gene is located on the Chr17. 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)

Rab5a RAB5A, member RAS oncogene family [Mus musculus (house mouse)]

Gene ID: 271457, updated on 7-Apr-2019

Summary



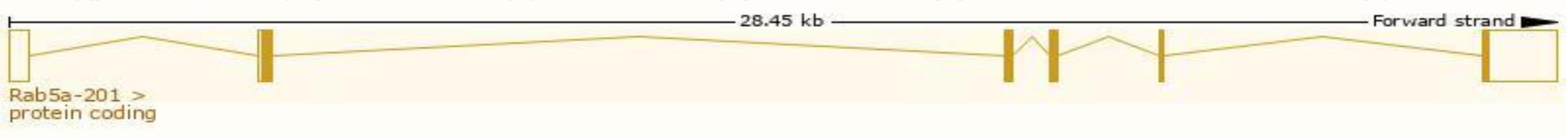
Official Symbol	Rab5a provided by MGI
Official Full Name	RAB5A, member RAS oncogene family provided by MGI
Primary source	MGI:MGI:105926
See related	Ensembl:ENSMUSG00000017831
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2410015H04Rik, A1663973, AU021172, nnyRab5a
Expression	Broad expression in CNS E11.5 (RPKM 27.6), CNS E18 (RPKM 24.0) and 25 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

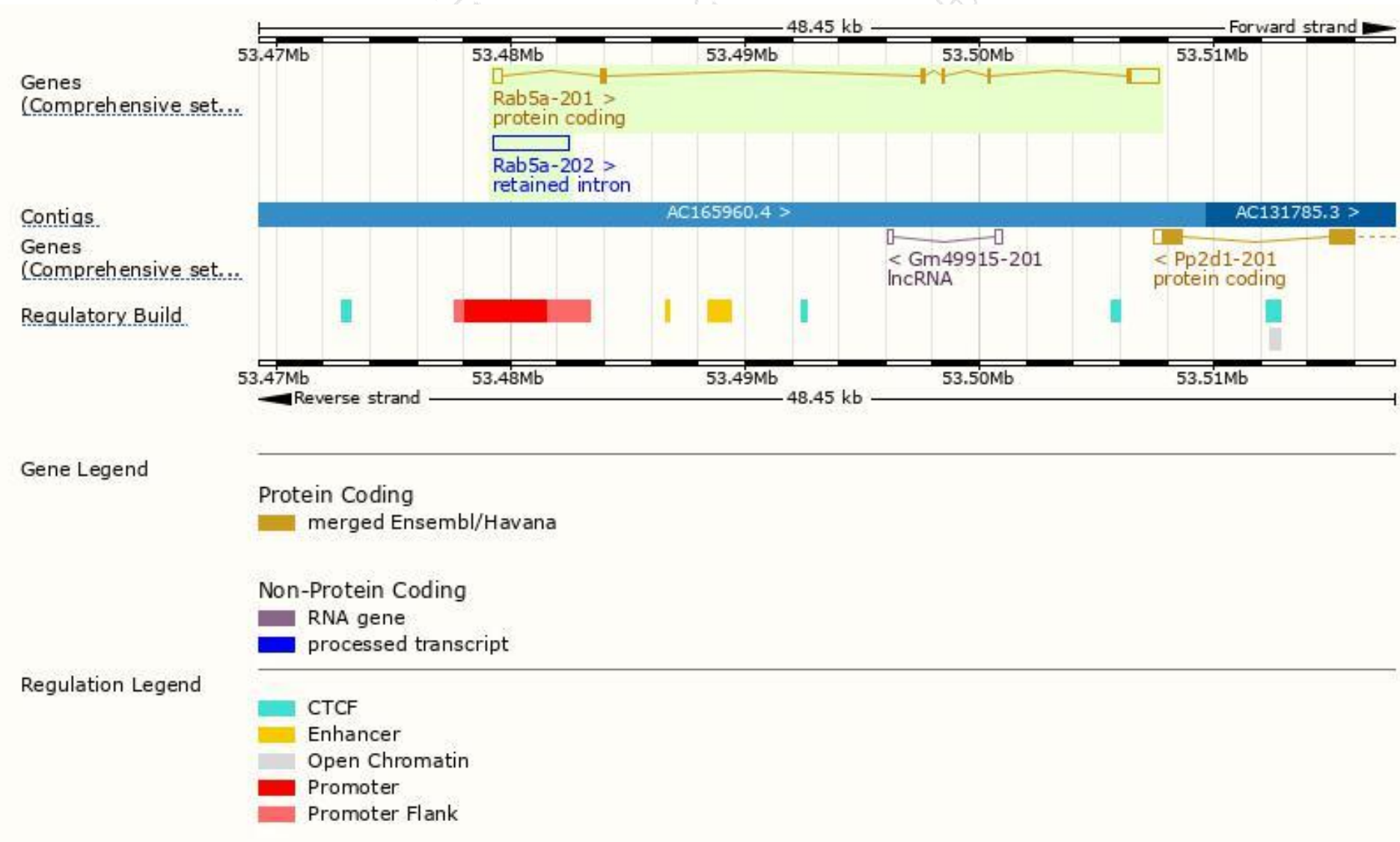
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rab5a-201	ENSMUST00000017975.6	2366	215aa	Protein coding	CCDS28879	Q9CQD1	TSL:1 GENCODE basic APPRIS P1
Rab5a-202	ENSMUST00000233347.1	3226	No protein	Retained intron	-	-	

The strategy is based on the design of *Rab5a-201* transcript,The transcription is shown below



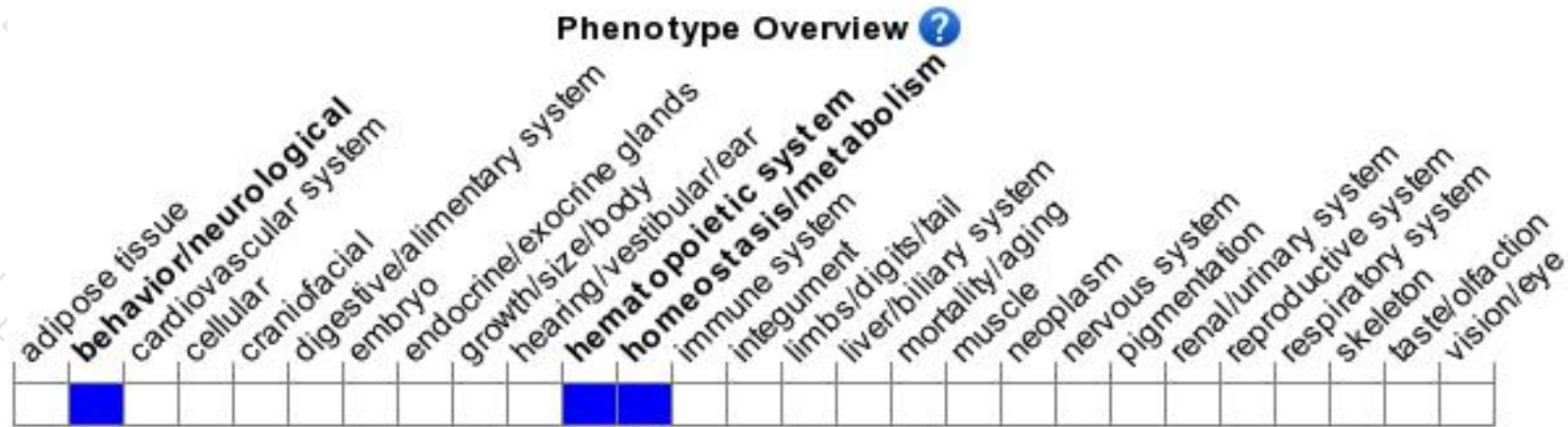
Genomic location distribution



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/>).

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

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