

***Rab10* Cas9-KO Strategy**

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Reviewer:

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Project Overview

Project Name

Rab10

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rab10* gene. The schematic diagram is as follows:



- The *Rab10* gene has 1 transcript. According to the structure of *Rab10* gene, exon2-exon5 of *Rab10-201* (ENSMUST00000021001.9) transcript is recommended as the knockout region. The region contains 392bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rab10* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit embryonic growth and development at E7.5 with reduced cell proliferation and complete resorption by E9.5.
- The *Rab10* gene is located on the Chr12. 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)

Rab10 RAB10, member RAS oncogene family [Mus musculus (house mouse)]

Gene ID: 19325, updated on 19-Mar-2019

Summary



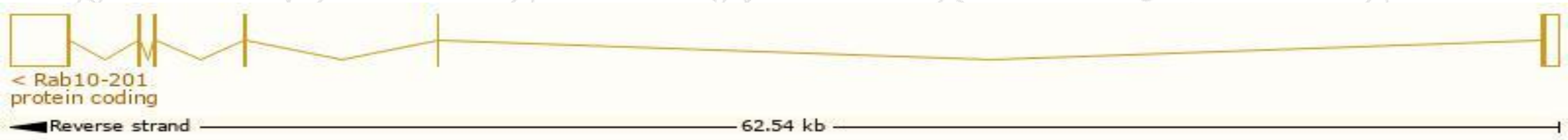
Official Symbol	Rab10 provided by MGI
Official Full Name	RAB10, member RAS oncogene family provided by MGI
Primary source	MGI:MGI:105066
See related	Ensembl:ENSMUSG00000020671
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AW107754
Expression	Ubiquitous expression in CNS E18 (RPKM 55.0), bladder adult (RPKM 47.9) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

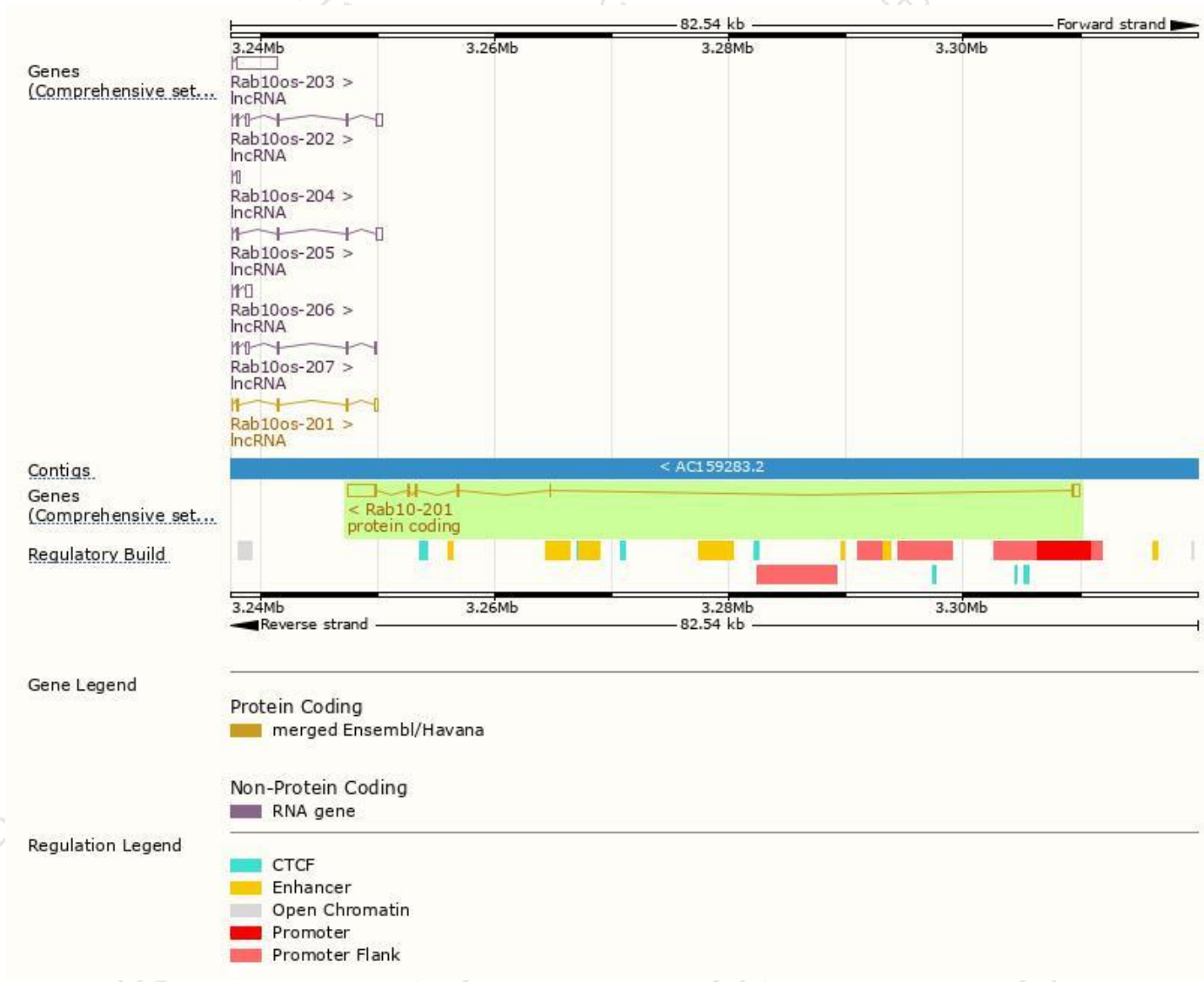
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rab10-201	ENSMUST00000021001.9	3513	200aa	Protein coding	CCDS25782	P61027 Q4FJL0	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Rab10-201* 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 embryonic growth and development at E7.5 with reduced cell proliferation and complete resorption by E9.5.

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

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