

Kcng4 Cas9-KO Strategy

Designer:

JiaYu

Reviewer:

Xiaojing Li

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Project Overview

Project Name

Kcng4

Project type

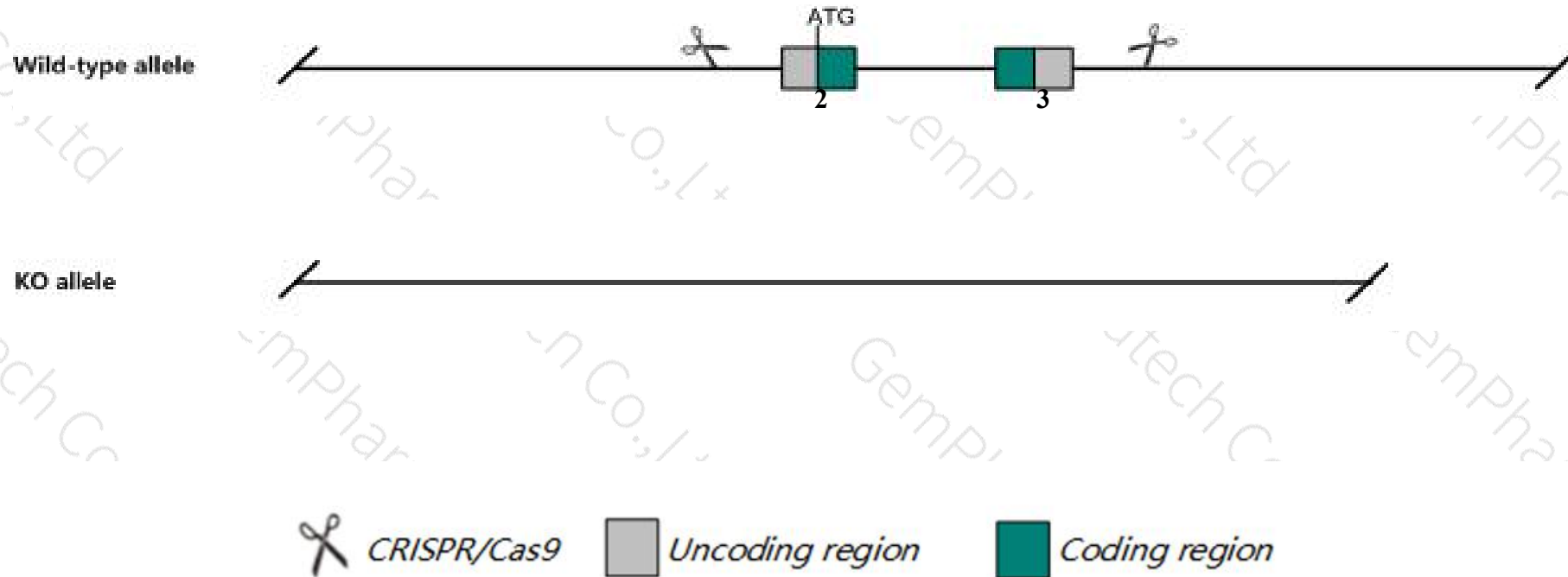
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Kcng4* gene. The schematic diagram is as follows:



- The *Kcng4* gene has 2 transcripts. According to the structure of *Kcng4* gene, exon2-exon3 of *Kcng4-201* (ENSMUST00000061828.9) transcript is recommended as the knockout region. The region contains all 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 *Kcng4* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Kcng4* gene is located on the Chr8. 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)

Kcng4 potassium voltage-gated channel, subfamily G, member 4 [Mus musculus (house mouse)]

Gene ID: 66733, updated on 31-Jan-2019

Summary



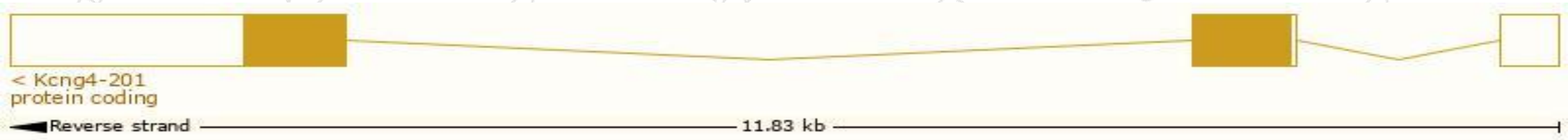
Official Symbol	Kcng4 provided by MGI
Official Full Name	potassium voltage-gated channel, subfamily G, member 4 provided by MGI
Primary source	MGI:MGI:1913983
See related	Ensembl:ENSMUSG00000045246
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4921535I01Rik, AW049024, KV6.3, KV6.4
Expression	Biased expression in cerebellum adult (RPKM 22.8), testis adult (RPKM 5.9) and 6 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

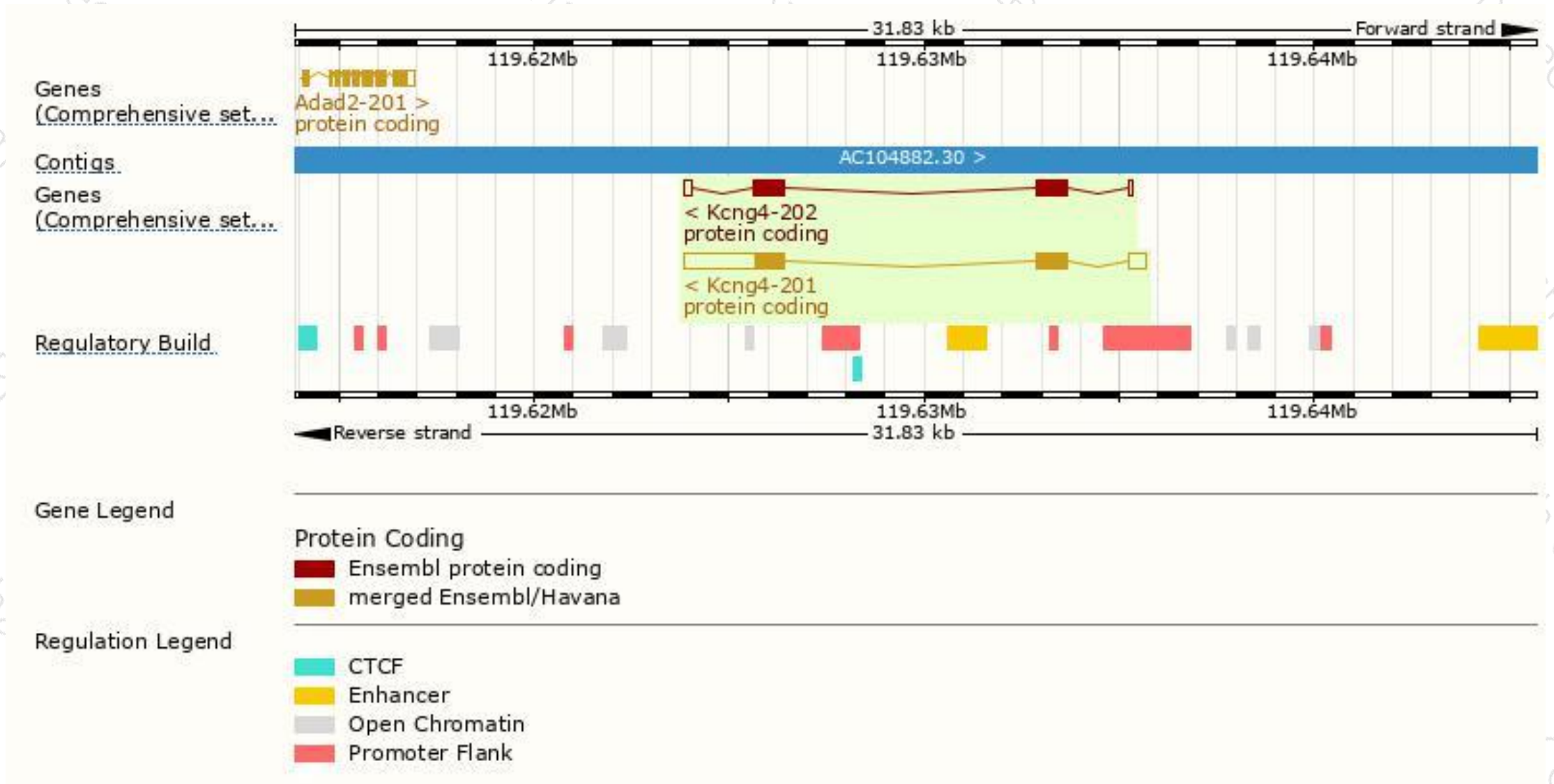
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kcng4-201	ENSMUST00000061828.9	3806	506aa	Protein coding	CCDS22710	Q80XM3	TSL:1 GENCODE basic APPRIS P1
Kcng4-202	ENSMUST00000164382.1	1875	506aa	Protein coding	CCDS22710	Q80XM3	TSL:5 GENCODE basic APPRIS P1

The strategy is based on the design of *Kcng4-201* transcript,The transcription is shown below



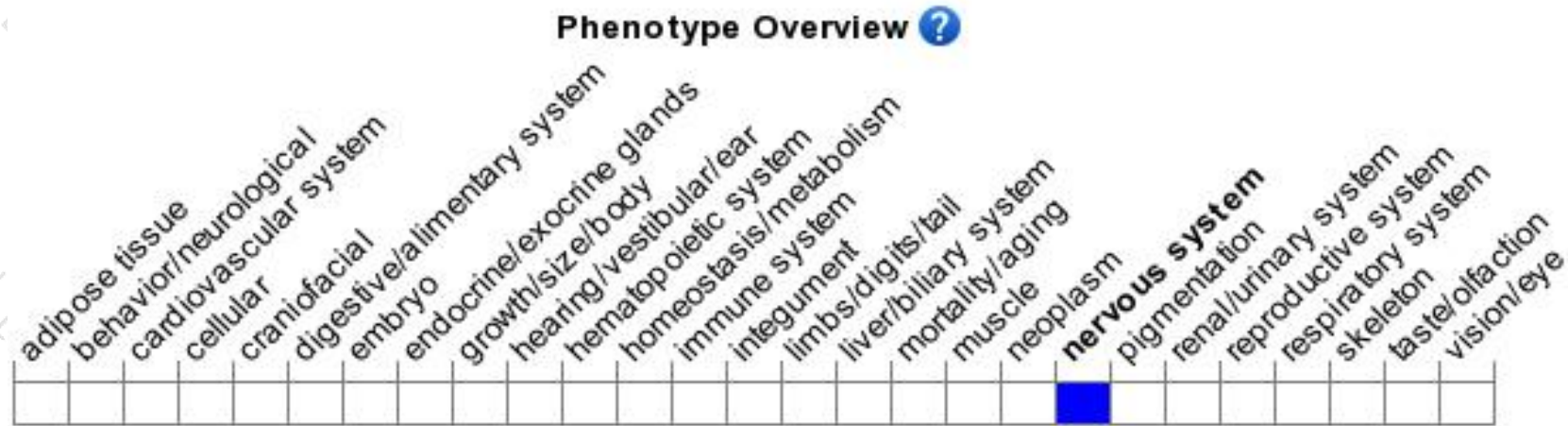
Genomic location distribution



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.

Tel: 400-9660890

