

Kcna6 Cas9-KO Strategy

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Date: 2020-02-19

Project Overview

Project Name

Kcna6

Project type

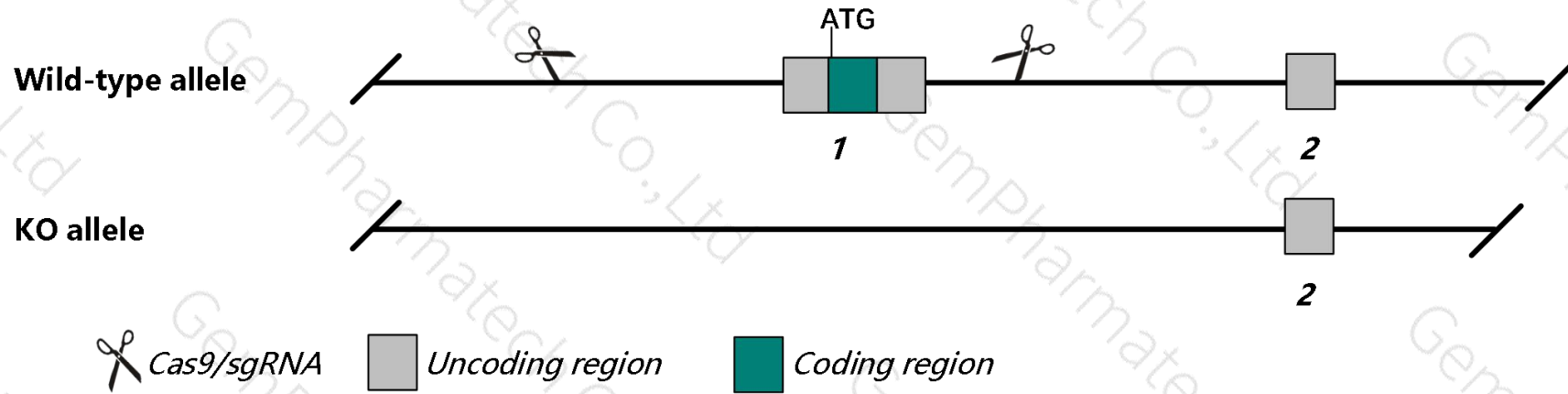
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Kcna6* gene. The schematic diagram is as follows:



- The *Kcna6* gene has 4 transcripts. According to the structure of *Kcna6* gene, exon1 of *Kcna6-204* (ENSMUST00000185333.1) 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 *Kcna6* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous mutation of this gene results in an increased thermal nociceptive threshold and in females an increase in circulating triglyceride levels.
- *Gm26673* gene will be deleted.
- The *Kcna6* gene is located on the Chr6. 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)

Kcna6 potassium voltage-gated channel, shaker-related, subfamily, member 6 [*Mus musculus* (house mouse)]

Gene ID: 16494, updated on 12-Aug-2019

Summary

- Official Symbol** Kcna6 provided by [MGI](#)
- Official Full Name** potassium voltage-gated channel, shaker-related, subfamily, member 6 provided by [MGI](#)
- Primary source** [MGI:MGI:96663](#)
- See related** [Ensembl:ENSMUSG00000038077](#)
- 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** Kv1.6; MK1.6
- Expression** Biased expression in cerebellum adult (RPKM 9.9), cortex adult (RPKM 8.5) and 7 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: 6 F3; 6 61.72 cM See Kcna6 in [Genome Data Viewer](#)

Exon count: 4

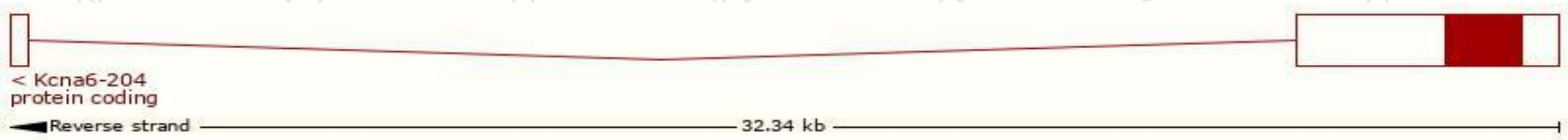
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	6	NC_000072.6 (126708328..126740834, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	6	NC_000072.5 (126658347..126690692, complement)

Transcript information (Ensembl)

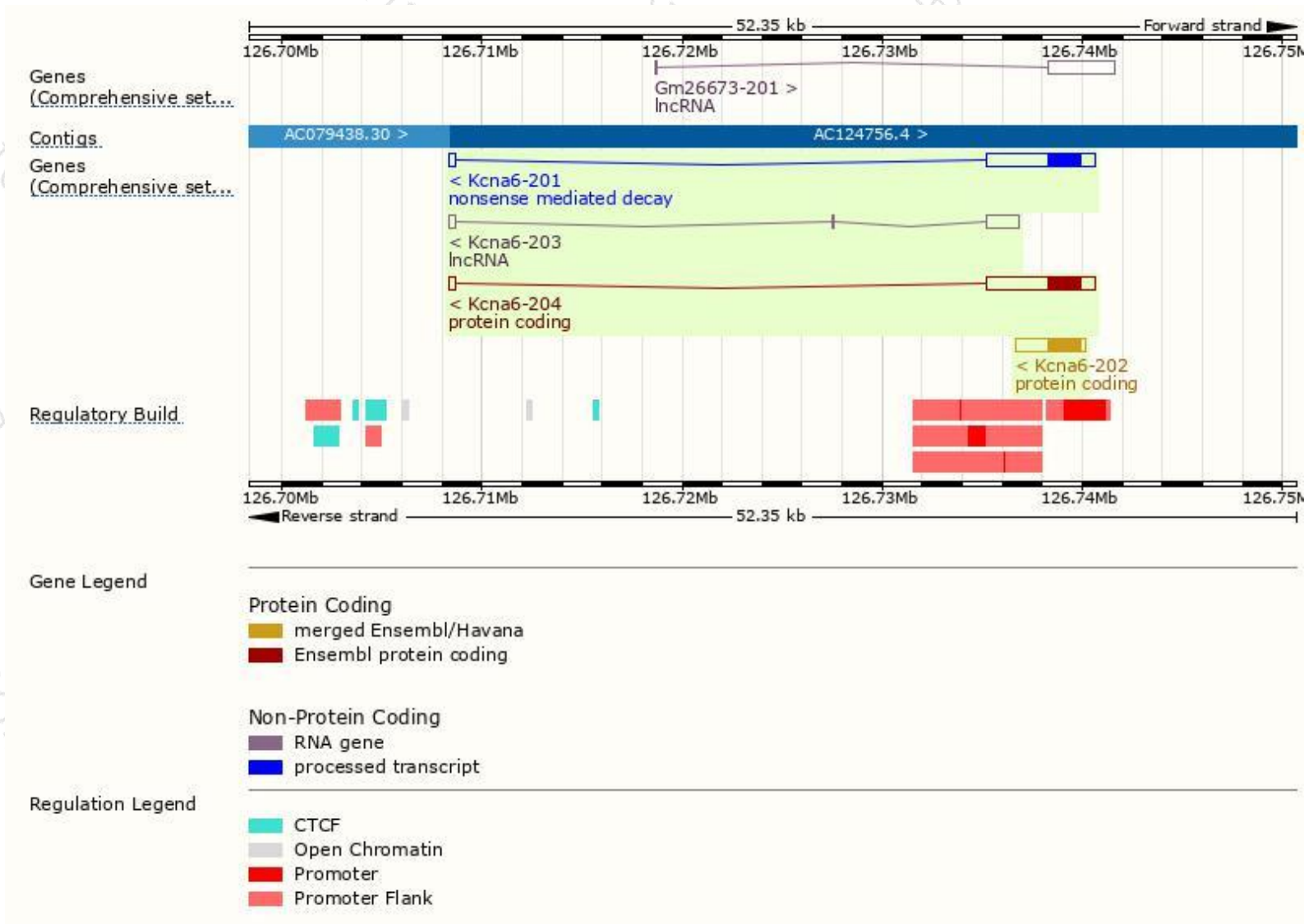
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kcna6-204	ENSMUST00000185333.1	5836	529aa	Protein coding	CCDS20556	Q61923	TSL:1 GENCODE basic APPRIS P1
Kcna6-202	ENSMUST00000112242.1	3464	529aa	Protein coding	CCDS20556	Q61923	TSL:NA GENCODE basic APPRIS P1
Kcna6-201	ENSMUST00000040751.5	5827	529aa	Nonsense mediated decay	CCDS20556	Q61923	TSL:1
Kcna6-203	ENSMUST00000128137.1	2080	No protein	lncRNA	-	-	TSL:1

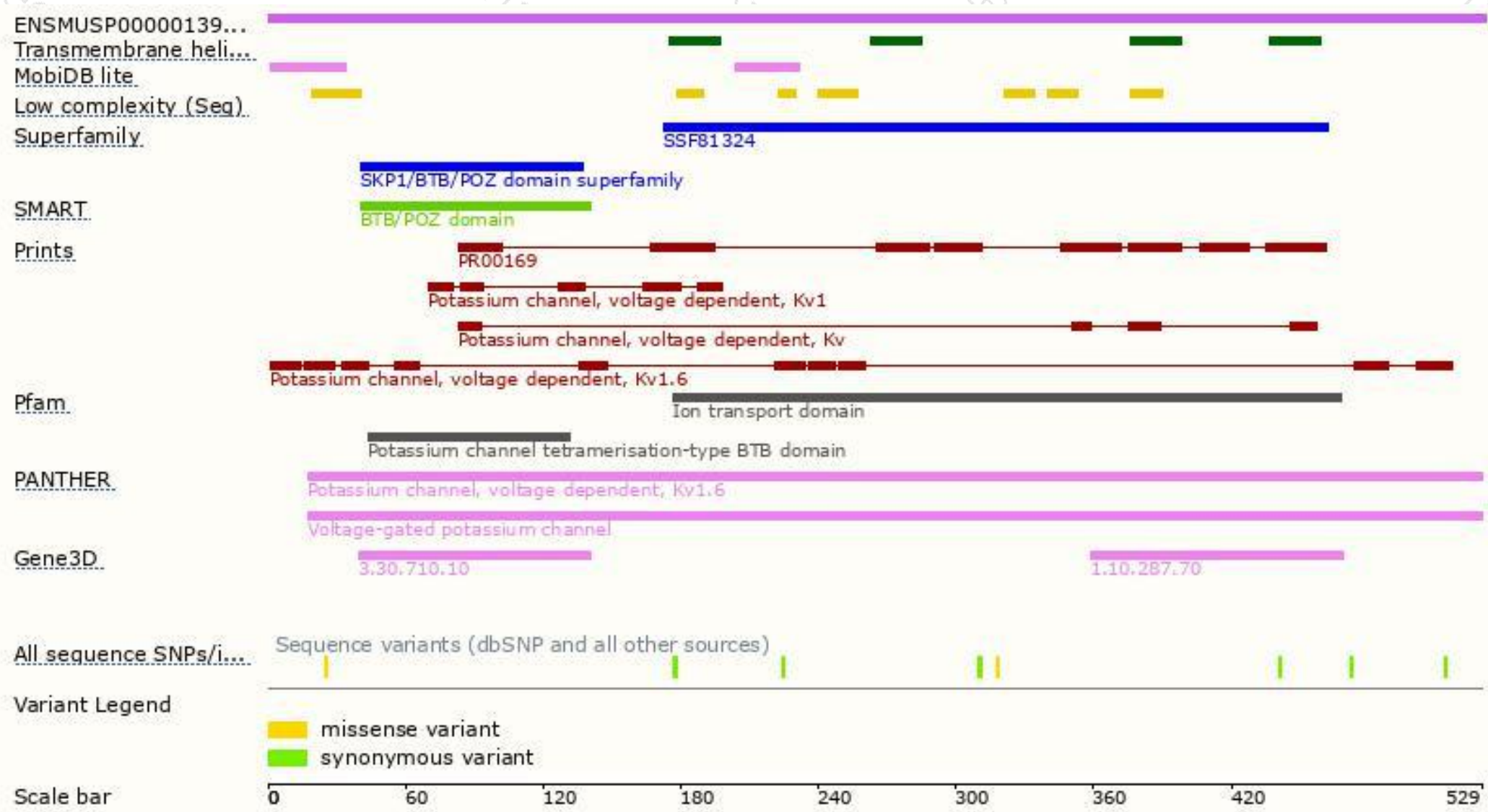
The strategy is based on the design of *Kcna6-204* transcript,The transcription is shown below



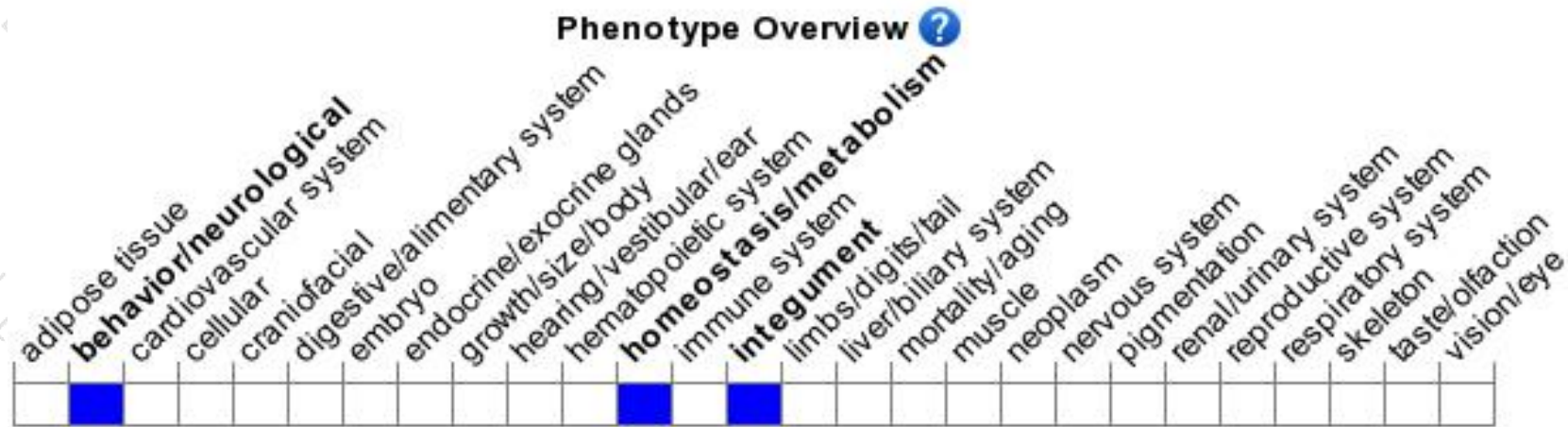
Genomic location distribution



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, Homozygous mutation of this gene results in an increased thermal nociceptive threshold and in females an increase in circulating triglyceride levels.

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

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