

Cacng4 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Cacng4

Project type

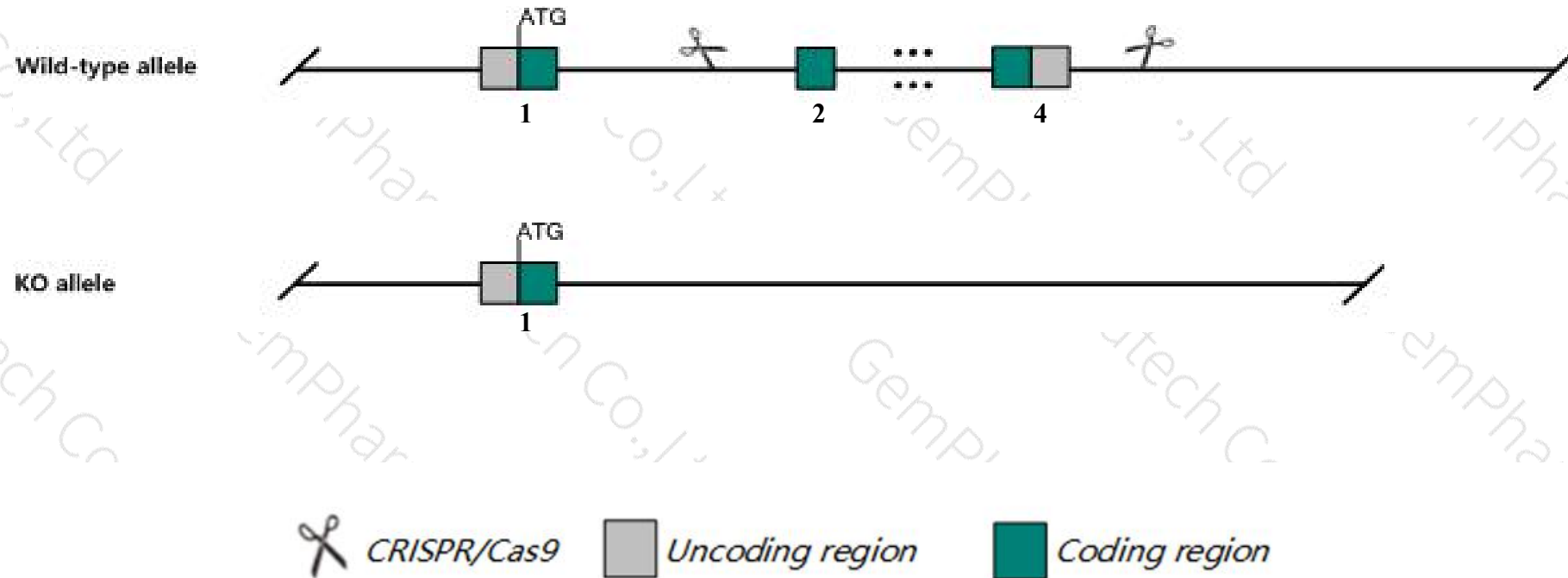
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cacng4* gene. The schematic diagram is as follows:



- The *Cacng4* gene has 2 transcripts. According to the structure of *Cacng4* gene, exon2-exon4 of *Cacng4-201* (ENSMUST00000021066.3) transcript is recommended as the knockout region. The region contains 764bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cacng4* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous mutant mice are viable, fertile and phenotypically normal with no ataxic gait or absence seizures.
- The *Cacng4* gene is located on the Chr11. 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)

Cacng4 calcium channel, voltage-dependent, gamma subunit 4 [Mus musculus (house mouse)]

Gene ID: 54377, updated on 19-Mar-2019

Summary



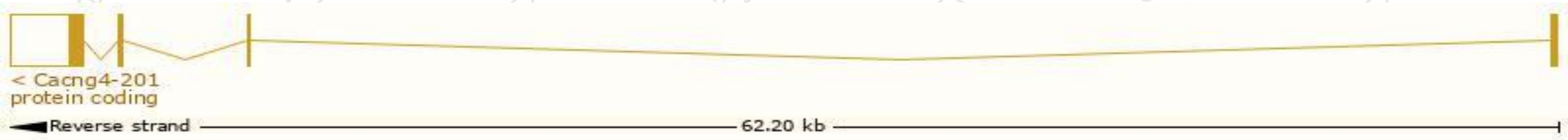
Official Symbol	Cacng4 provided by MGI
Official Full Name	calcium channel, voltage-dependent, gamma subunit 4 provided by MGI
Primary source	MGI:MGI:1859167
See related	Ensembl:ENSMUSG00000020723
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI413107, AW491861
Expression	Biased expression in whole brain E14.5 (RPKM 47.6), CNS E18 (RPKM 44.4) and 7 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

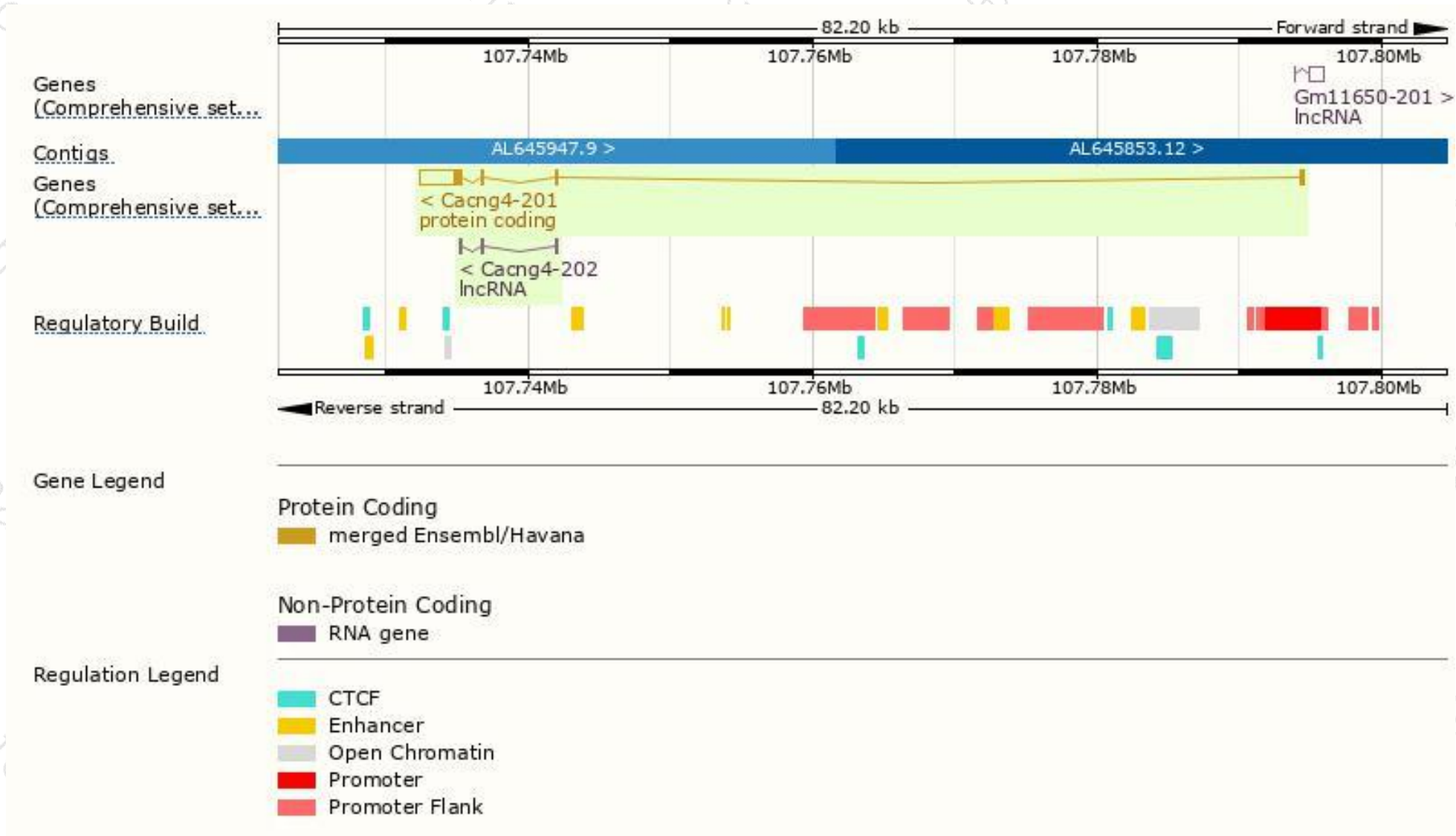
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cacng4-201	ENSMUST00000021066.3	3500	327aa	Protein coding	CCDS25571	A2AAU2 Q9JJV4	TSL:1 GENCODE basic APPRIS P1
Cacng4-202	ENSMUST00000134076.1	422	No protein	lncRNA	-	-	TSL:2

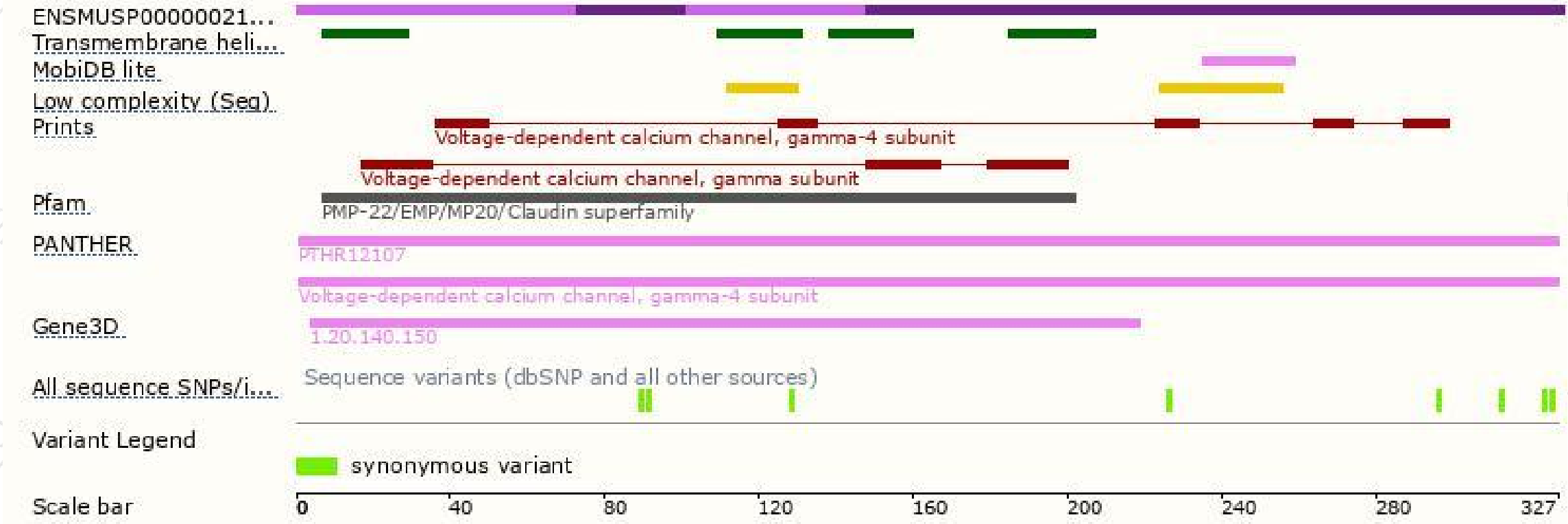
The strategy is based on the design of *Cacng4-201* transcript,The transcription is shown below



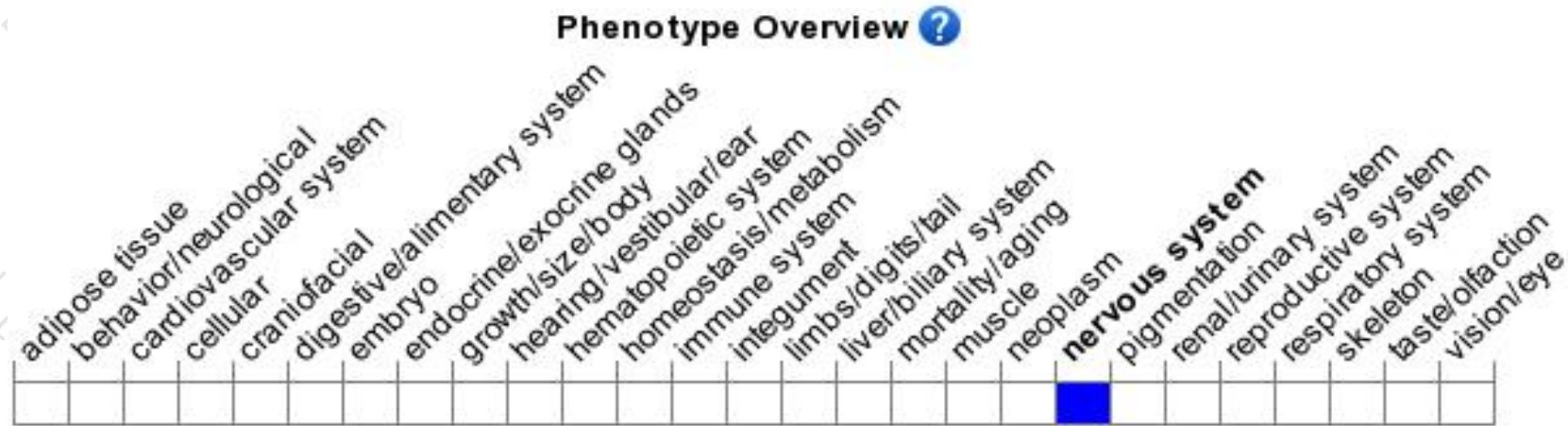
Genomic location distribution



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, Homozygous mutant mice are viable, fertile and phenotypically normal with no ataxic gait or absence seizures.

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

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