

Calb1 Cas9-KO Strategy

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Project Overview

Project Name

Calb1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Calb1* gene. The schematic diagram is as follows:



- The *Calb1* gene has 3 transcripts. According to the structure of *Calb1* gene, exon3-exon11 of *Calb1-201* (ENSMUST00000029876.1) transcript is recommended as the knockout region. The region contains most 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 *Calb1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous targeted mutants show severely impairment in motor coordination and Purkinje cells in the cerebellum show changes of synaptically evoked postsynaptic calcium transients.
- The *Calb1* gene is located on the Chr4. 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)

Calb1 calbindin 1 [*Mus musculus* (house mouse)]

Gene ID: 12307, updated on 15-Oct-2019

Summary

- Official Symbol

Calb1 provided by MGI
- Official Full Name

calbindin 1 provided by MGI
- Primary source

MGI:MGI:88248
- See related

Ensembl:ENSMUSG00000028222
- 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

CB; Calb; Calb-1; Brain-2
- Expression

Biased expression in cerebellum adult (RPKM 80.7), kidney adult (RPKM 24.9) and 5 other tissues [See more](#)
- Orthologs

[human](#) [all](#)

Genomic context

Location:

4 A2; 4 6.66 cM

See Calb1 in [Genome Data Viewer](#)

Exon count:

11

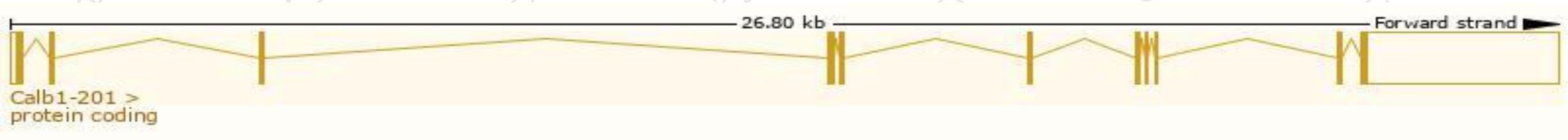
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	4	NC_000070.6 (15881264..15906709)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	4	NC_000070.5 (15808411..15833856)

Transcript information (Ensembl)

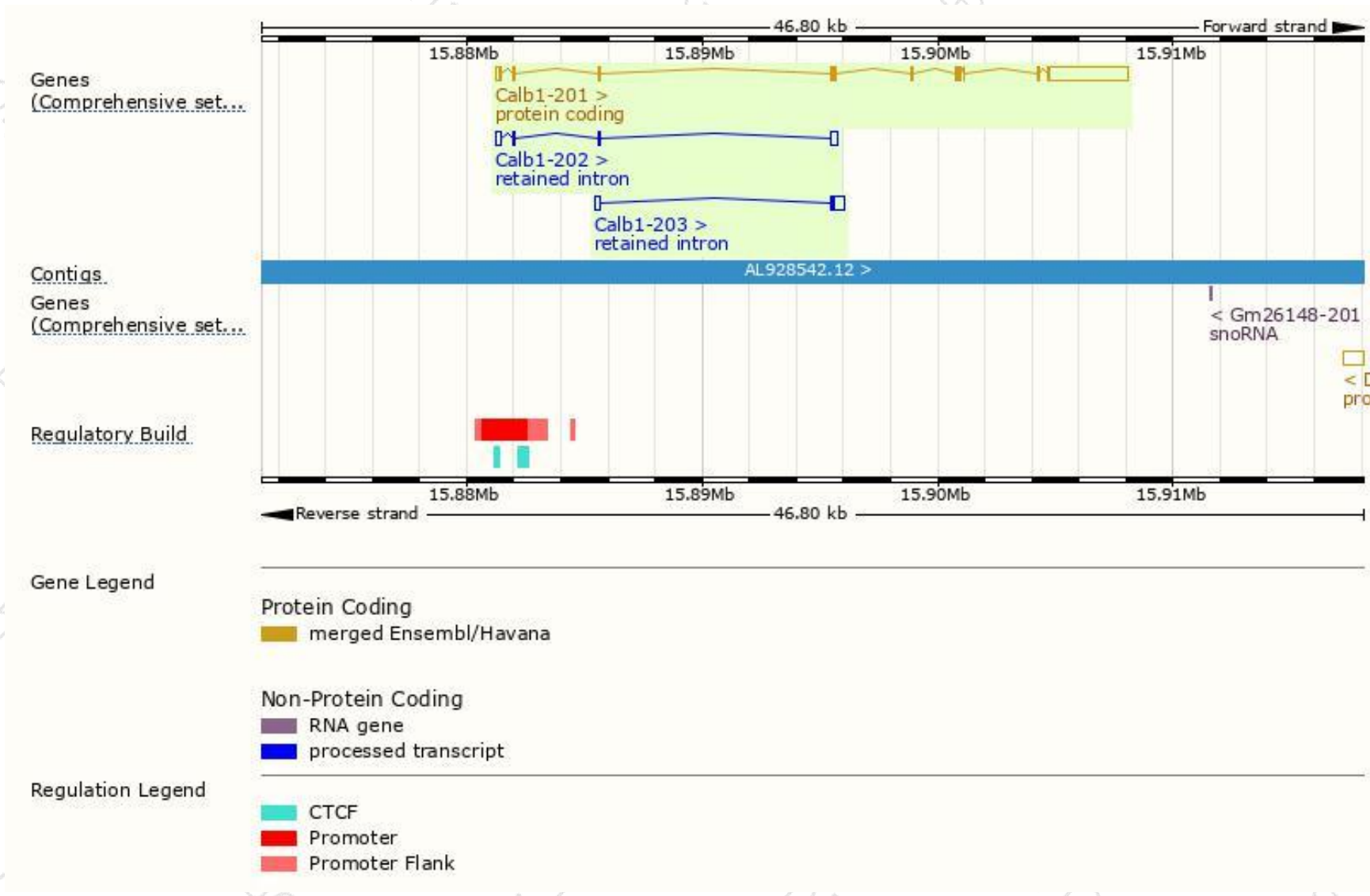
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Calb1-201	ENSMUST00000029876.1	4222	261aa	Protein coding	CCDS17984	P12658	TSL:1 GENCODE basic APPRIS P1
Calb1-203	ENSMUST00000141336.1	736	No protein	Retained intron	-	-	TSL:3
Calb1-202	ENSMUST00000136266.1	701	No protein	Retained intron	-	-	TSL:2

The strategy is based on the design of *Calb1-201* transcript,The transcription is shown below



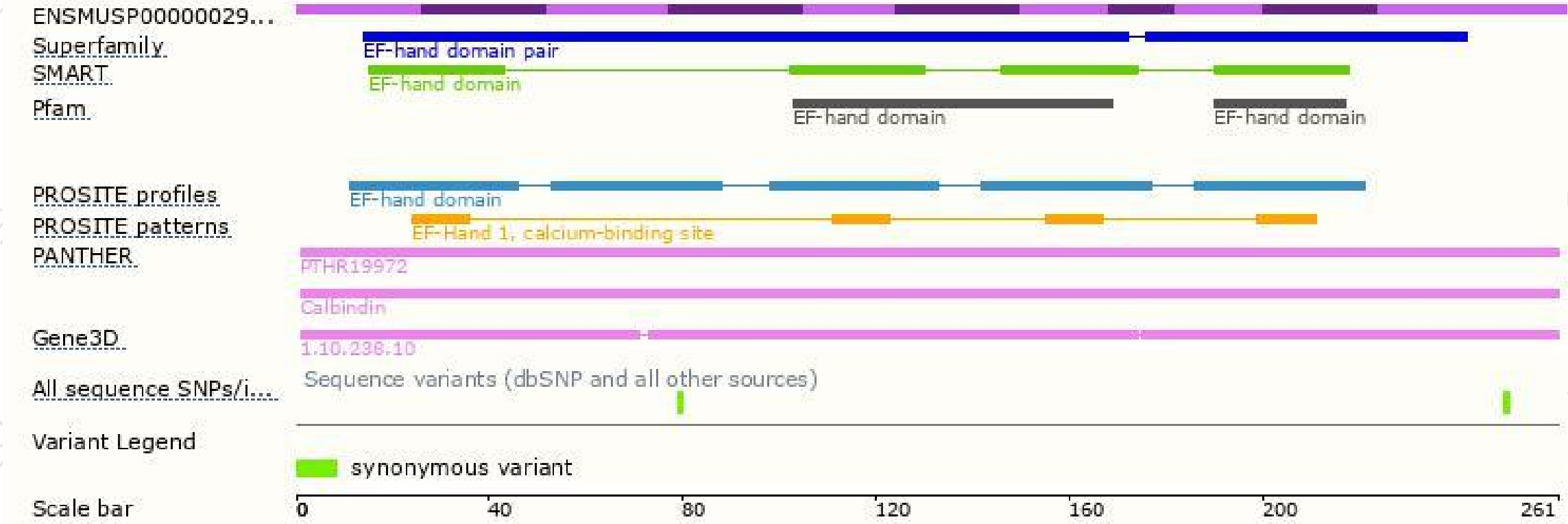
Genomic location distribution



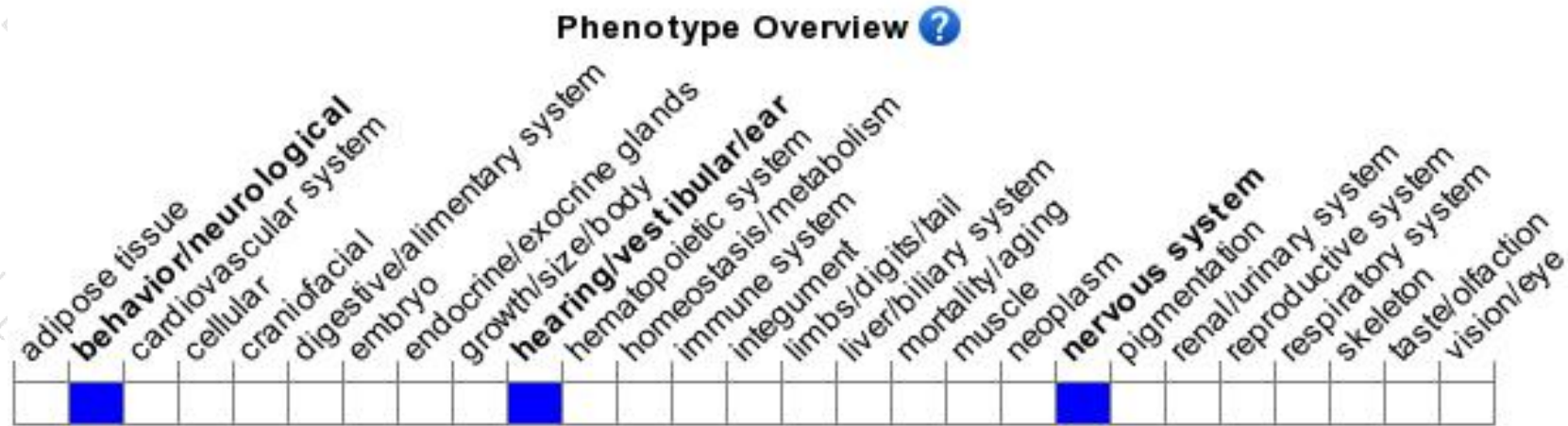
Protein domain



集萃药康
GemPharmatech



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 targeted mutants show severely impairment in motor coordination and Purkinje cells in the cerebellum show changes of synaptically evoked postsynaptic calcium transients.

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

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