

Gca Cas9-KO Strategy

Designer:

Daohua Xu

Reviewer :

Huimin Su

Design Date:

2020-3-23

Project Overview

Project Name

Gca

Project type

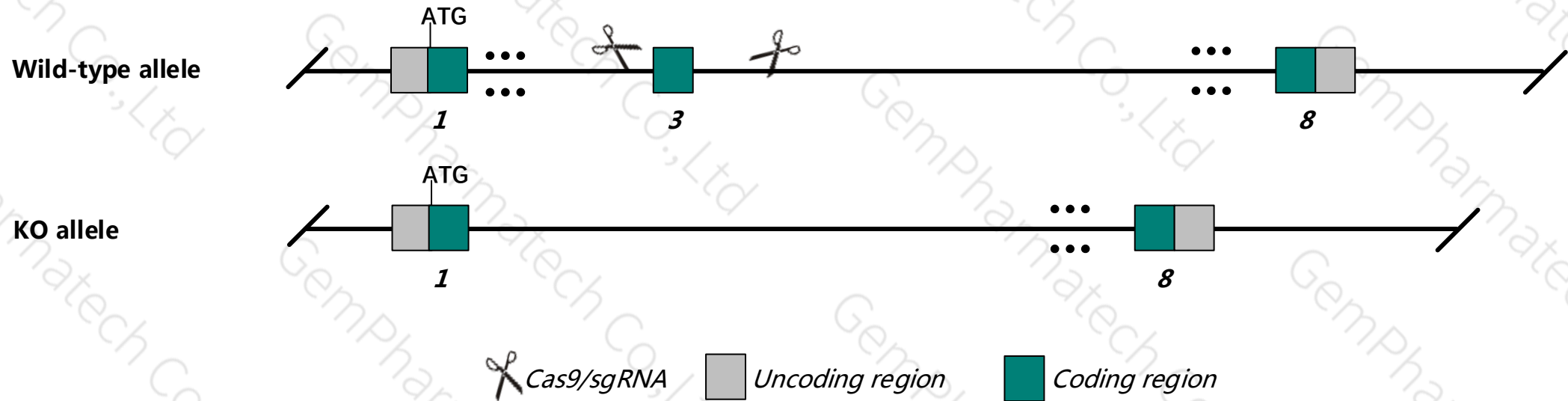
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



Technical routes

- The *Gca* gene has 2 transcripts. According to the structure of *Gca* gene, exon3 of *Gca*-201 (ENSMUST00000028257.2) transcript is recommended as the knockout region. The region contains 70bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gca* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9, sgRNA Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

- According to the existing MGI data , Mice homozygous for disruptions in this gene are essentially normal. However they do demonstrate an increased resistance to endotoxic shock.
- The *Gca* gene is located on the Chr2. 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)

Gca grancalcin [*Mus musculus* (house mouse)]

Gene ID: 227960, updated on 13-Mar-2020

Summary

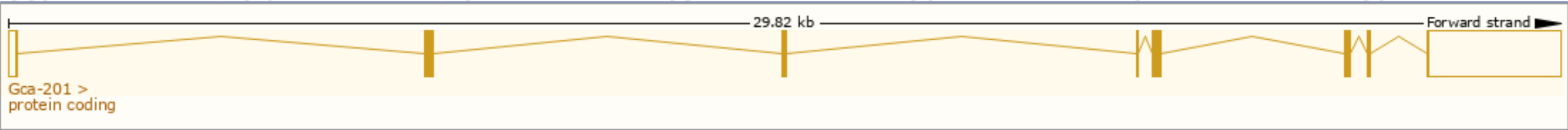
Official Symbol	Gca provided by MGI
Official Full Name	grancalcin provided by MGI
Primary source	MGI:MGI:1918521
See related	Ensembl:ENSMUSG00000026893
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI573844; 5133401E04Rik
Expression	Ubiquitous expression in cortex adult (RPKM 6.2), frontal lobe adult (RPKM 5.7) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

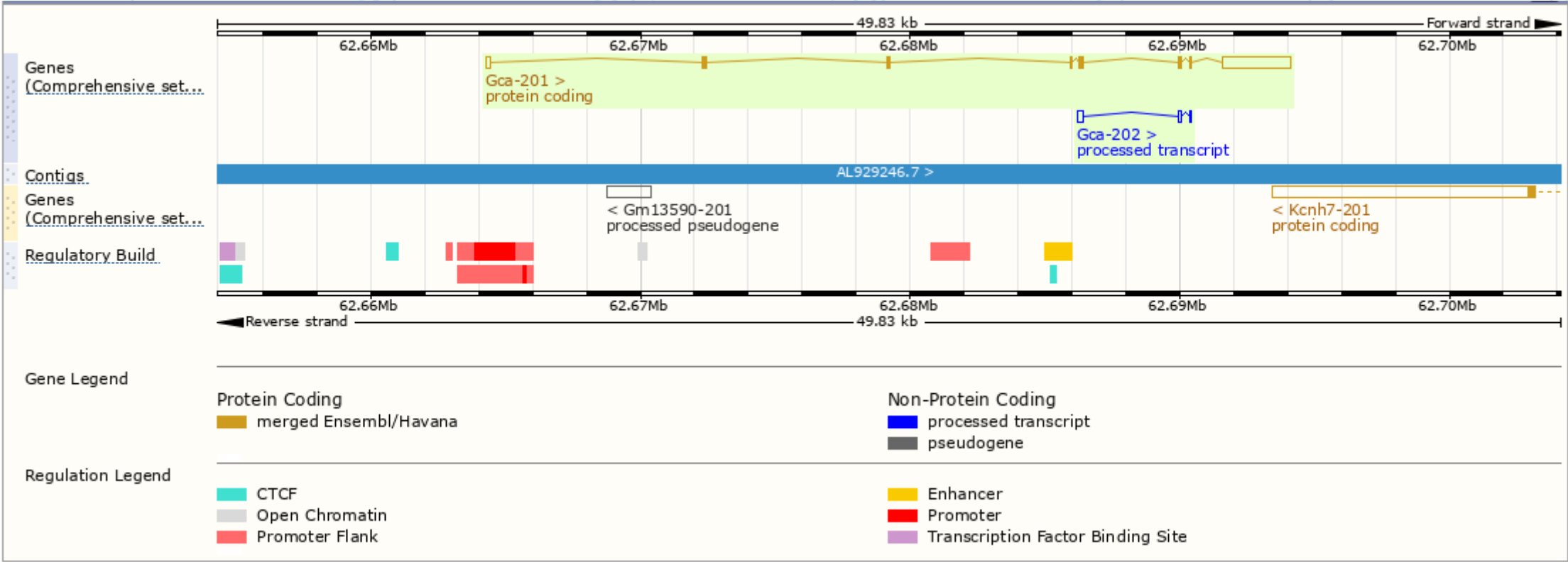
The gene has 2 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gca-201	ENSMUST00000028257.2	3330	220aa	Protein coding	CCDS16069	Q8VC88	TSL:1 GENCODE basic APPRIS P1
Gca-202	ENSMUST00000148083.1	382	No protein	Processed transcript	-	-	TSL:2

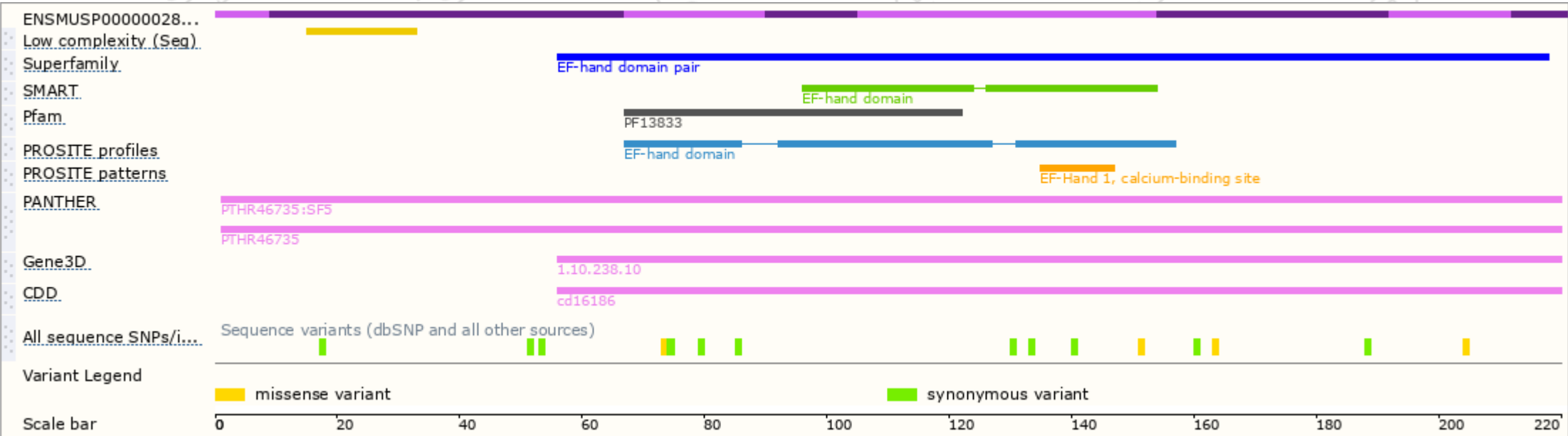
The strategy is based on the design of *Gca-201* transcript, The transcription is shown below



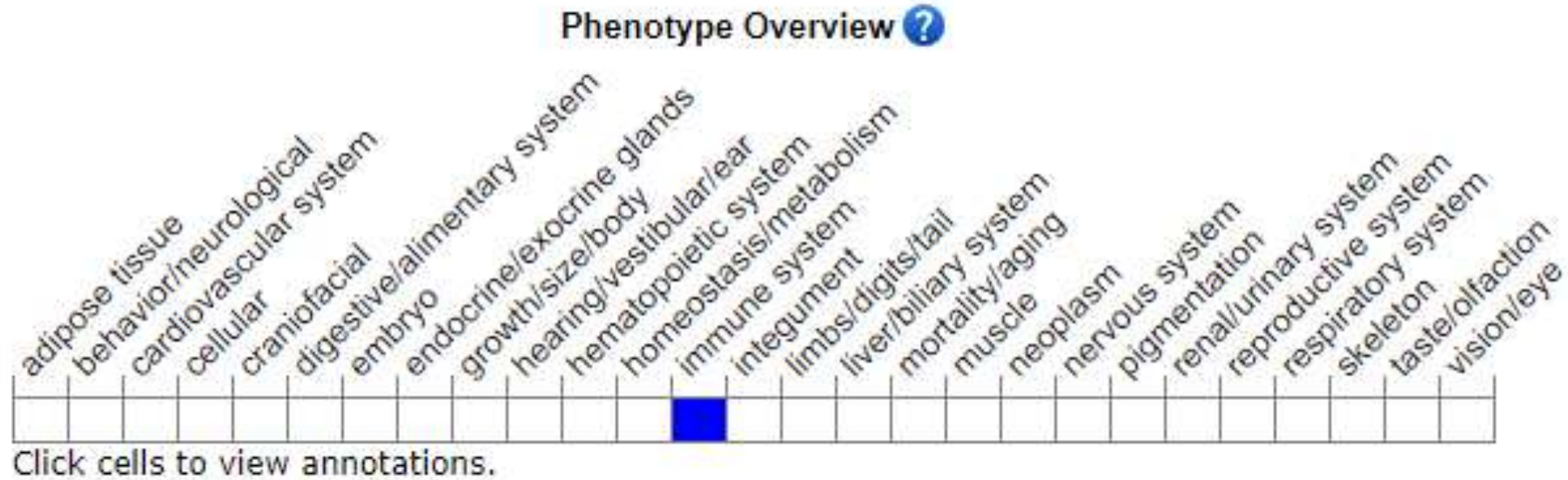
Genomic location distribution



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 disruptions in this gene are essentially normal. However they do demonstrate an increased resistance to endotoxic shock.

If you have any questions, you are welcome to inquire.
Tel: 025-5864 1534



集萃药康生物科技
GemPharmatech Co.,Ltd

