

***Gng4* Cas9-CKO Strategy**

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Reviewer:

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Design Date:

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Project Overview

Project Name

Gng4

Project type

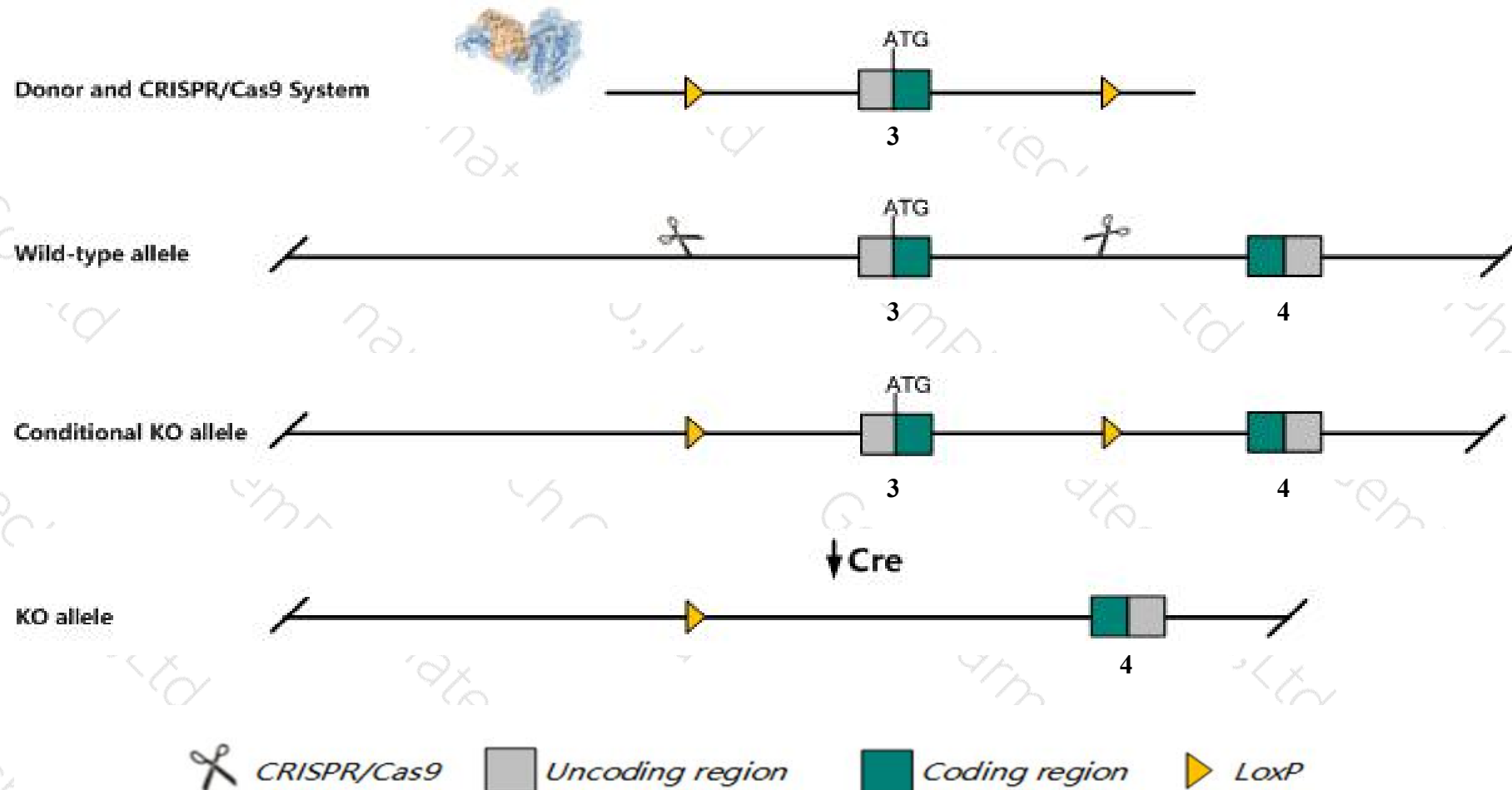
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gng4* gene. The schematic diagram is as follows:



- The *Gng4* gene has 3 transcripts. According to the structure of *Gng4* gene, exon3 of *Gng4-201* (ENSMUST00000021734.8) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gng4* gene. The brief process is as follows: CRISPR/Cas9 system and 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- The *Gng4* gene is located on the Chr13. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Gng4 guanine nucleotide binding protein (G protein), gamma 4 [Mus musculus (house mouse)]

Gene ID: 14706, updated on 31-Jan-2019

Summary



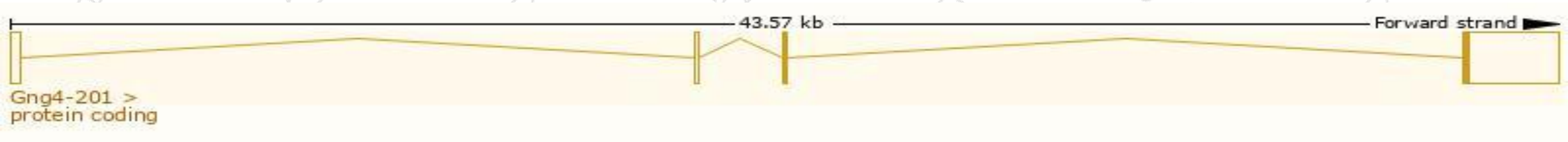
Official Symbol	Gng4 provided by MGI
Official Full Name	guanine nucleotide binding protein (G protein), gamma 4 provided by MGI
Primary source	MGI:MGI:102703
See related	Ensembl:ENSMUSG000000021303
Gene type	protein coding
RefSeq status	REVIEWED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Summary	This gene encodes the gamma subunit of the heterotrimeric G-proteins that are comprised of alpha, beta and gamma subunits. Upon activation by G protein-coupled receptors, the beta-gamma heterodimer dissociates from the alpha subunit to activate downstream signaling events. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Dec 2014]
Expression	Biased expression in frontal lobe adult (RPKM 69.2), CNS E18 (RPKM 22.4) and 3 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

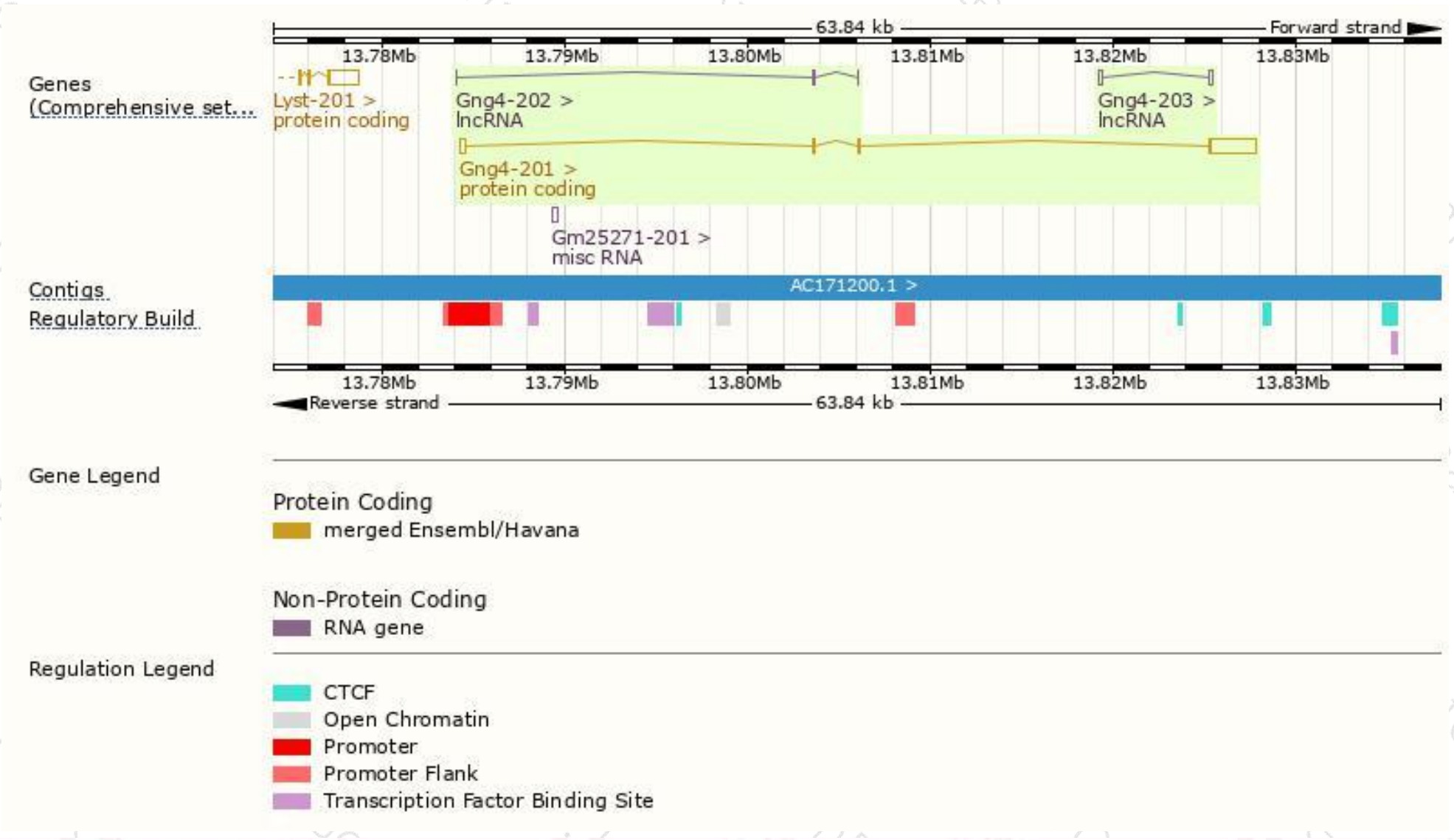
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gng4-201	ENSMUST00000021734.8	3179	75aa	Protein coding	CCDS36598	P50153	TSL:1 GENCODE basic APPRIS P1
Gng4-203	ENSMUST00000220727.1	324	No protein	lncRNA	-	-	TSL:3
Gng4-202	ENSMUST00000138783.7	206	No protein	lncRNA	-	-	TSL:3

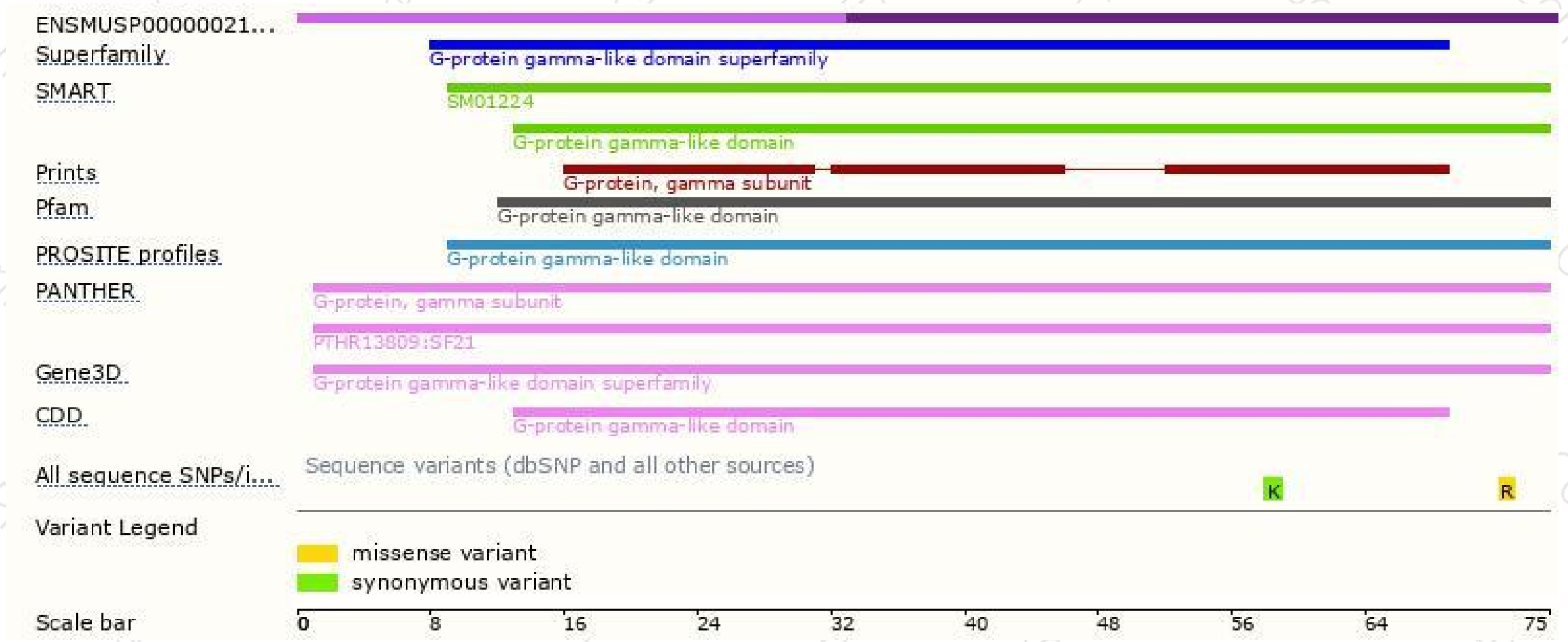
The strategy is based on the design of *Gng4-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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