

Chrnb2 Cas9-CKO Strategy Rohalana Koch Co.

JiaYu

Project Overview



Project Name

Chrnb2

Project type

Cas9-CKO

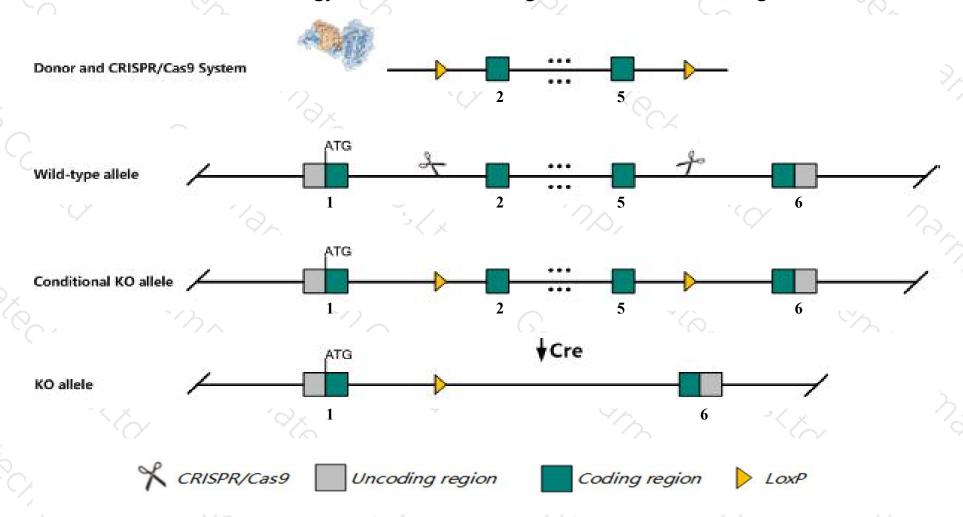
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Chrnb2 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Chrnb2* gene has 3 transcripts. According to the structure of *Chrnb2* gene, exon2-exon5 of *Chrnb2-201* (ENSMUST00000029562.4) transcript is recommended as the knockout region. The region contains 1271bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Chrnb2* 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



- ➤ According to the existing MGI data, Homozygotes for targeted null mutations have impaired responses to nicotine, but show improved passive avoidance behavior. With age, mutants show more neurodegeneration and alterations of the visual system, with decreased cortical visual acuity.
- > The *Chrnb2* gene is located on the Chr3. 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)



Chrnb2 cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal) [Mus musculus (house mouse)]

Gene ID: 11444, updated on 19-Feb-2019

Summary

☆ ?

Official Symbol Chrnb2 provided by MGI

Official Full Name cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal) provided by MGI

Primary source MGI:MGI:87891

See related Ensembl:ENSMUSG00000027950

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 Acrb-2, Acrb2, C030030P04Rik, [b]2-nAchR

Expression Biased expression in whole brain E14.5 (RPKM 17.9), cerebellum adult (RPKM 14.9) and 8 other tissuesSee more

Orthologs <u>human all</u>

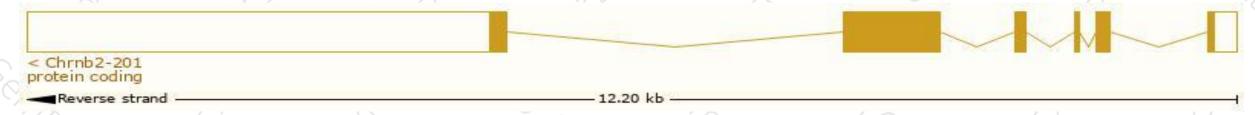
Transcript information (Ensembl)



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

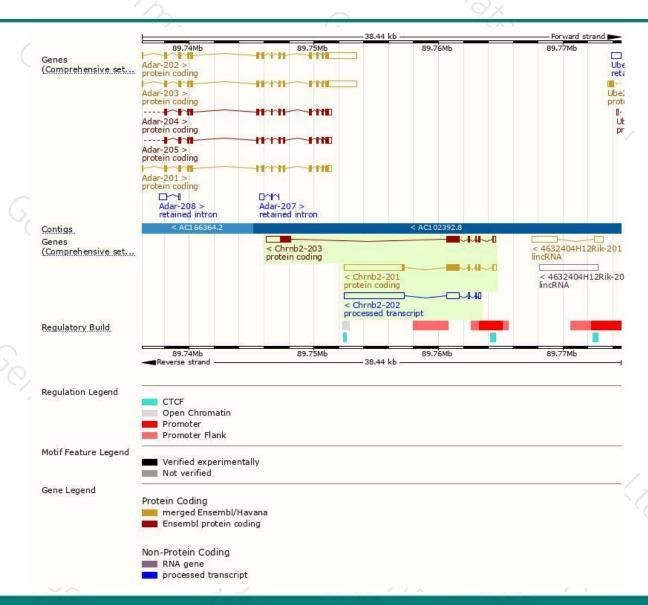
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Chrnb2-201	ENSMUST00000029562.4	6399	501aa	Protein coding	CCDS17515	Q9ERK7	TSL:1 GENCODE basic APPRIS P1
Chrnb2-203	ENSMUST00000200558.4	3482	727aa	Protein coding	-	A0A0G2JG64	TSL:1 GENCODE basic
Chrnb2-202	ENSMUST00000199372.1	6145	No protein	Processed transcript	020	12	TSL:5

The strategy is based on the design of Chrnb2-201 transcript, The transcription is shown below



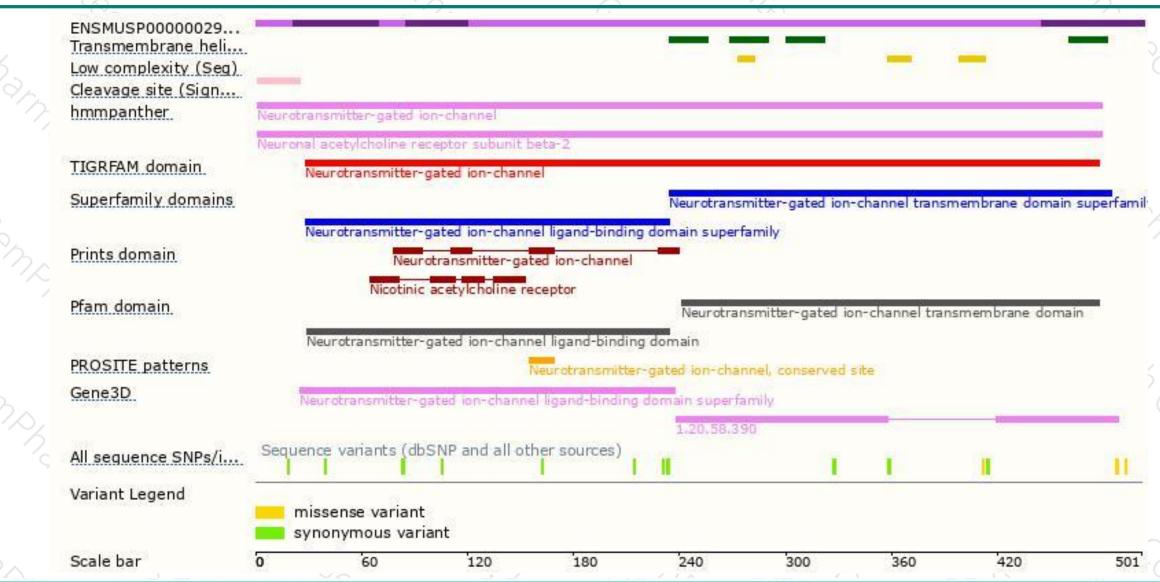
Genomic location distribution





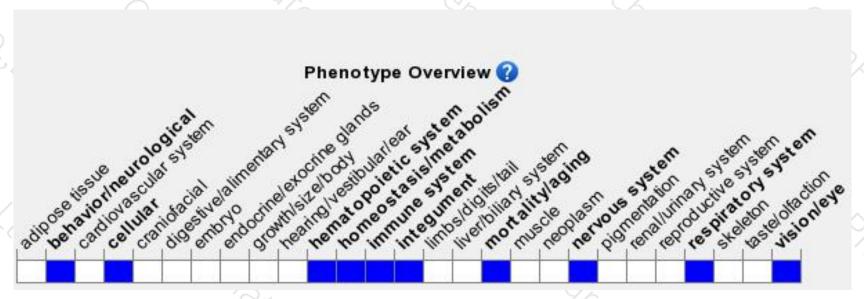
Protein domain





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, Homozygotes for targeted null mutations have impaired responses to nicotine, but show improved passive avoidance behavior. With age, mutants show more neurodegeneration and alterations of the visual system, with decreased cortical visual acuity.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





