

Chrna7 Cas9-CKO Strategy

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Project Overview



Project Name

Chrna7

Project type

Cas9-CKO

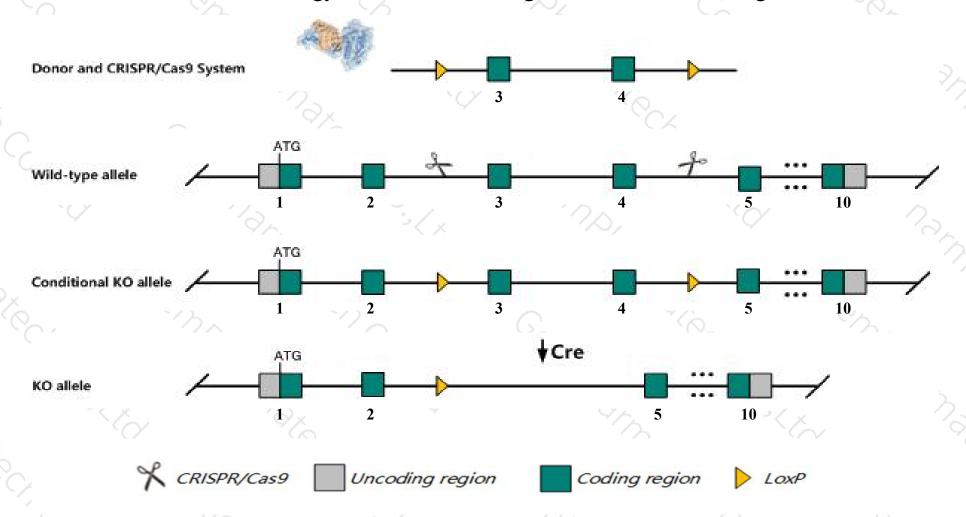
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Chrna7* gene. The schematic diagram is as follows:



Technical routes



- The *Chrna7* gene has 2 transcripts. According to the structure of *Chrna7* gene, exon3-exon4 of *Chrna7-201* (ENSMUST00000032738.6) transcript is recommended as the knockout region. The region contains 155bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Chrna7* 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, Nullizygous mice lack hippocampal fast nicotinic currents but show nicotine-induced seizures as well as altered anxiety behavior, fertility defects, airway basal cell hyperplasia. and higher TNF sythesis when endotoxemic. Newborns homozygous for a knock-in allele die with increased neuron apoptosis.
- > The *Chrna7* gene is located on the Chr7. 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)



Chrna7 cholinergic receptor, nicotinic, alpha polypeptide 7 [Mus musculus (house mouse)]

Gene ID: 11441, updated on 12-Mar-2019

Summary

↑ ?

Official Symbol Chrna7 provided by MGI

Official Full Name cholinergic receptor, nicotinic, alpha polypeptide 7 provided by MGI

Primary source MGI:MGI:99779

See related Ensembl:ENSMUSG00000030525

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 Acra7, alpha7, nAchR

Expression Biased expression in adrenal adult (RPKM 6.7), CNS E18 (RPKM 3.8) and 10 other tissuesSee more

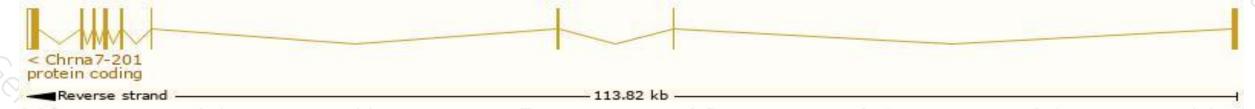
Transcript information (Ensembl)



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

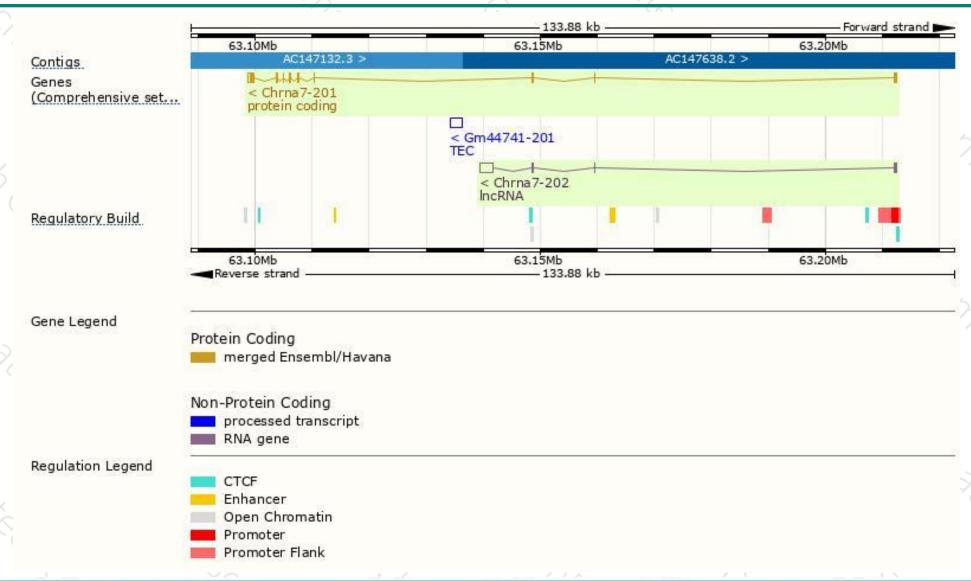
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Chrna7-201	ENSMUST00000032738.6	2078	502aa	Protein coding	CCDS21329	P49582 Q53YJ9	TSL:1 GENCODE basic APPRIS P1
Chrna7-202	ENSMUST00000208304.1	2814	No protein	Processed transcript			TSL:1

The strategy is based on the design of Chrna7-201 transcript, The transcription is shown below



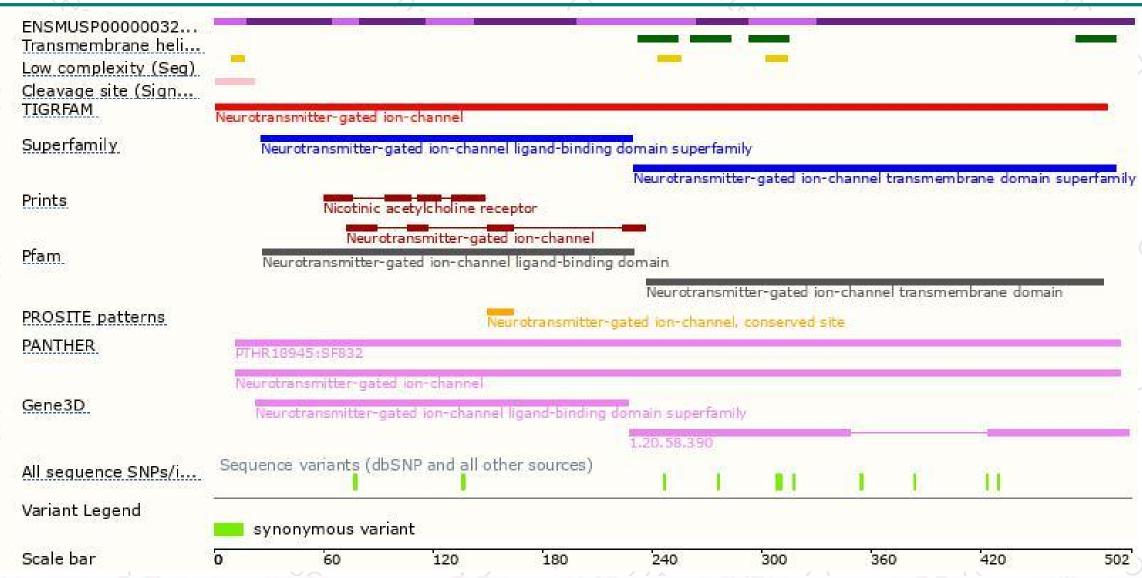
Genomic location distribution





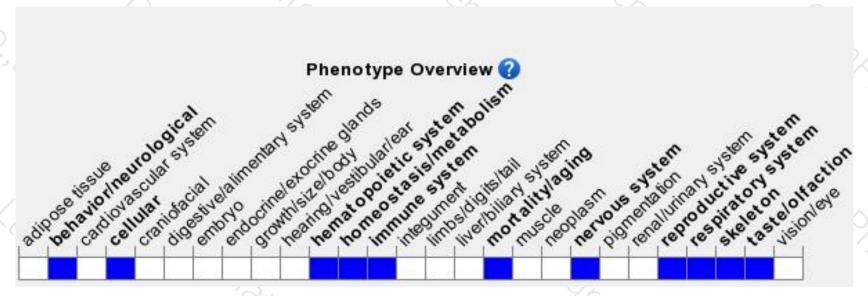
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

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





