

Clic5 Cas9-CKO Strategy

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Reviewer:

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Design Date:

2019-9-10

Project Overview



Project Name Clic5

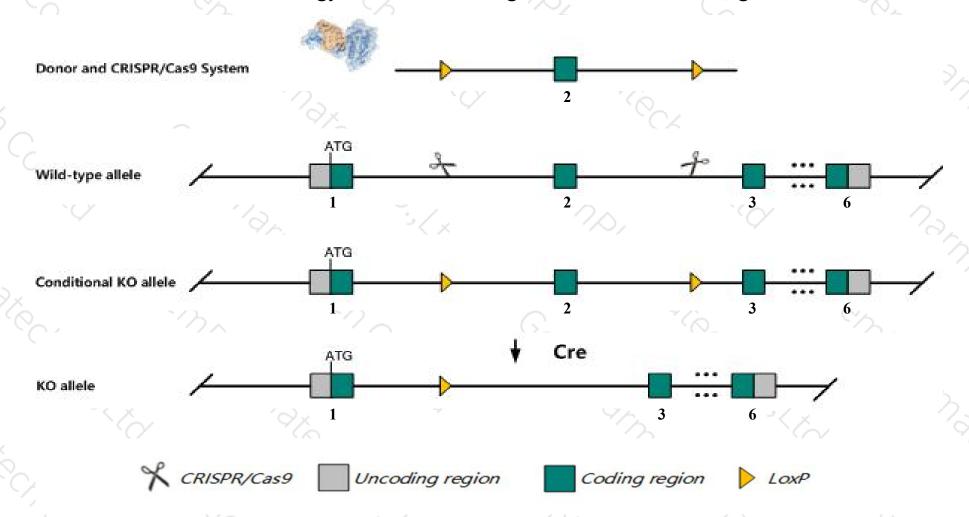
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Clic5 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Clic5* gene has 2 transcripts. According to the structure of *Clic5* gene, exon2 of *Clic5-201*(ENSMUST00000024755.6) transcript is recommended as the knockout region. The region contains 110bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Clic5* 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, Mice homozygous for a spontaneous mutation exhibit head bobbing and circling behavior and complete deafness by 7-8 months of age caused by dysmorphic stereocilia and progressive hair cell degeneration. Mice are resistant to diet-induced obesity and are unable to maintain energy reserves.

 Cataracts develop with age. Renal glomerulus defects are present.
- The *Clic5* gene is located on the Chr17. 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)



Clic5 chloride intracellular channel 5 [Mus musculus (house mouse)]

Gene ID: 224796, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Clic5 provided by MGI

Official Full Name chloride intracellular channel 5 provided by MGI

Primary source MGI:MGI:1917912

See related Ensembl:ENSMUSG00000023959

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 5730531E12Rik, B330005L24, Gm322, jbg, nmf318

Expression Biased expression in lung adult (RPKM 40.1), large intestine adult (RPKM 14.4) and 10 other tissuesSee more

Orthologs human all

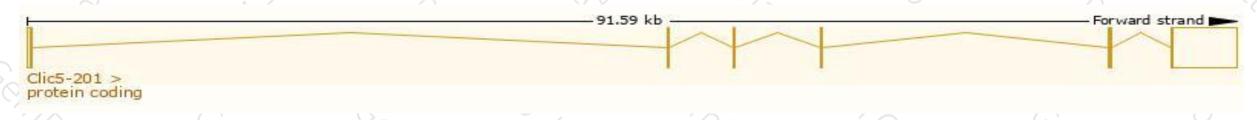
Transcript information (Ensembl)



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

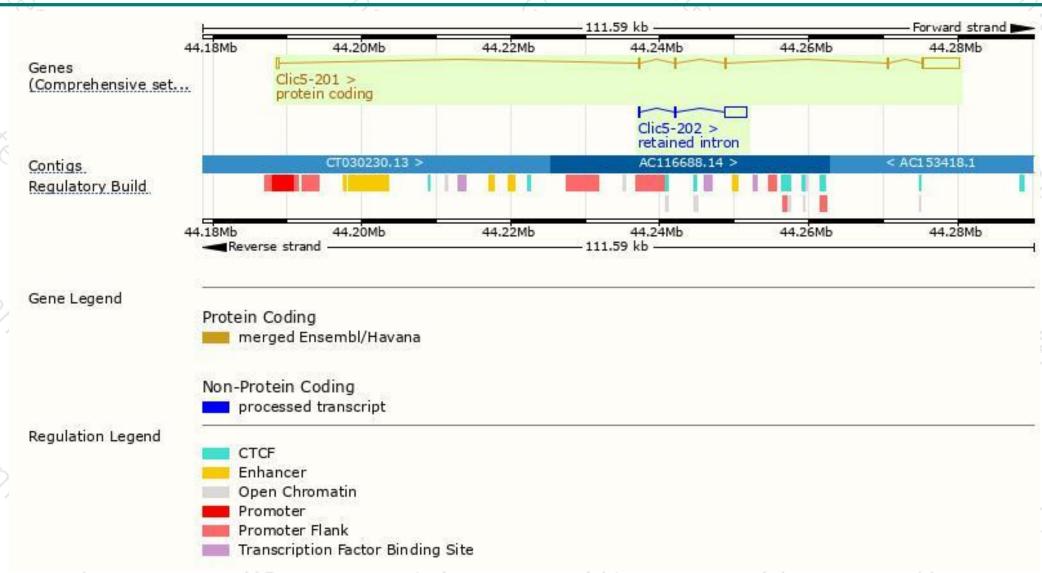
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Clic5-201	ENSMUST00000024755.6	5862	<u>251aa</u>	Protein coding	CCDS28804	Q8BXK9	TSL:1 GENCODE basic APPRIS P1
Clic5-202	ENSMUST00000233193.1	3263	No protein	Retained intron	1940	15 5	

The strategy is based on the design of Clic5-201 transcript, The transcription is shown below



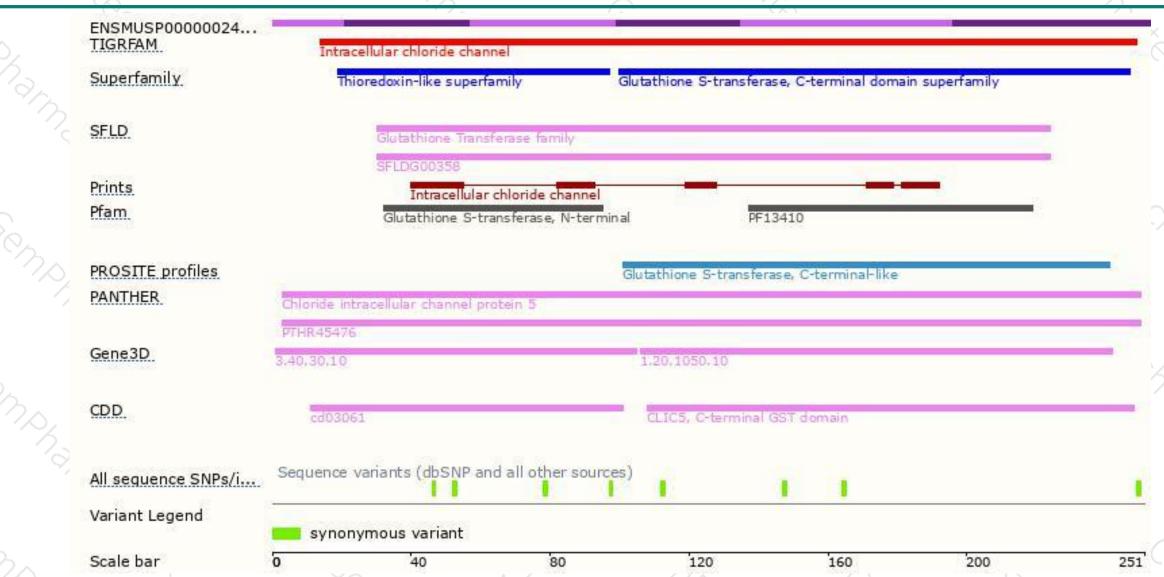
Genomic location distribution





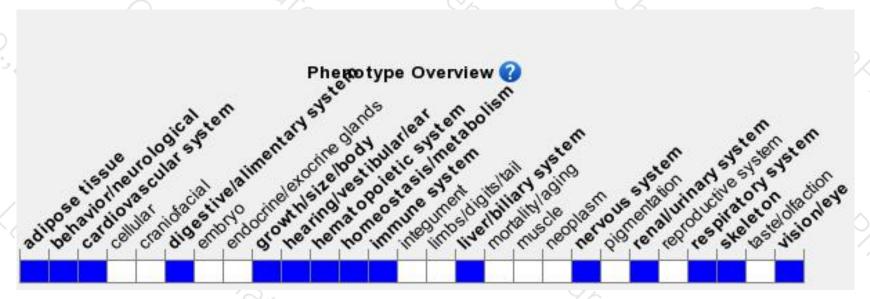
Protein domain





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





