

Cdhr3 Cas9-KO Strategy

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Project Overview

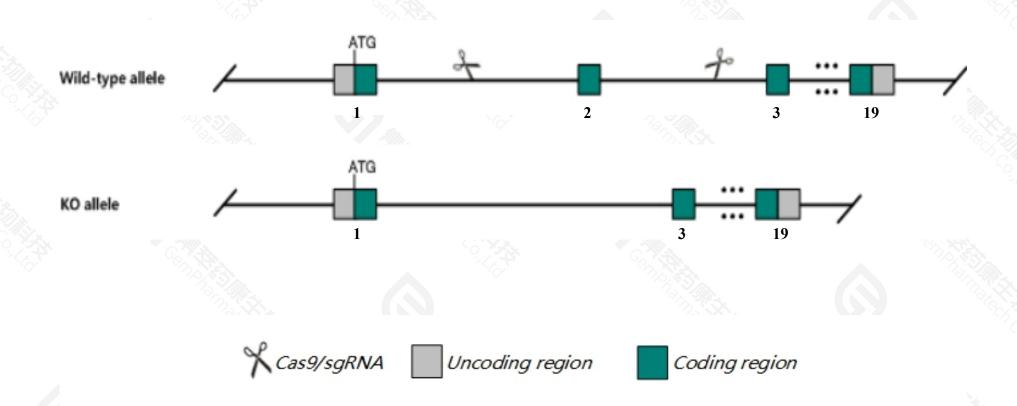


Project Name	Cdhr3
Project type	Cas9-KO
Strain background	C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Cdhr3* gene. The schematic diagram is as follows:



Technical routes



- > The *Cdhr3* gene has 4 transcripts. According to the structure of *Cdhr3* gene, exon2 of *Cdhr3*-201(ENSMUST00000095774.2) transcript is recommended as the knockout region. The region contains 203bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Cdhr3* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA 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.

Notice



- > The *Cdhr3* gene is located on the Chr12. 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)



Cdhr3 cadherin-related family member 3 [Mus musculus (house mouse)]

Gene ID: 68764, updated on 13-Mar-2020

Summary



Official Symbol Cdhr3 provided by MGI

Official Full Name cadherin-related family member 3 provided by MGI

Primary source MGI:MGI:1916014

See related Ensembl:ENSMUSG00000035860

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 1110049B09Rik, BB172694, Cdh28

Expression Biased expression in lung adult (RPKM 4.3), ovary adult (RPKM 1.8) and 4 other tissuesSee more

Orthologs <u>human</u> <u>all</u>

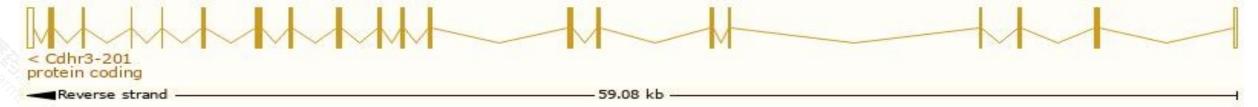
Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

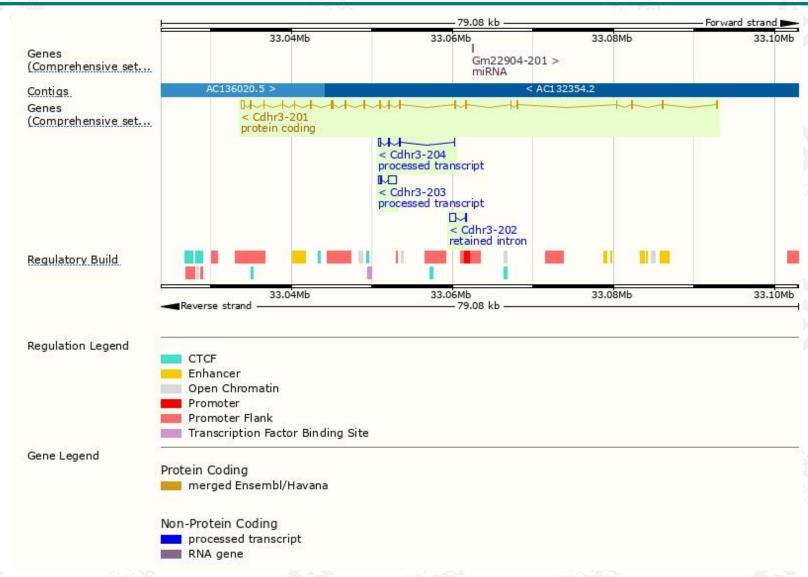
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cdhr3-201	ENSMUST00000095774.2	2953	<u>831aa</u>	Protein coding	CCDS36430	Q8BL00	TSL:1 GENCODE basic APPRIS P1
Cdhr3-203	ENSMUST00000219158.1	1209	No protein	Processed transcript	-	-	TSL:1
Cdhr3-204	ENSMUST00000219453.1	604	No protein	Processed transcript	-	-	TSL:3
Cdhr3-202	ENSMUST00000219065.1	881	No protein	Retained intron	-	-	TSL:1

The strategy is based on the design of *Cdhr3-201* transcript, the transcription is shown below:



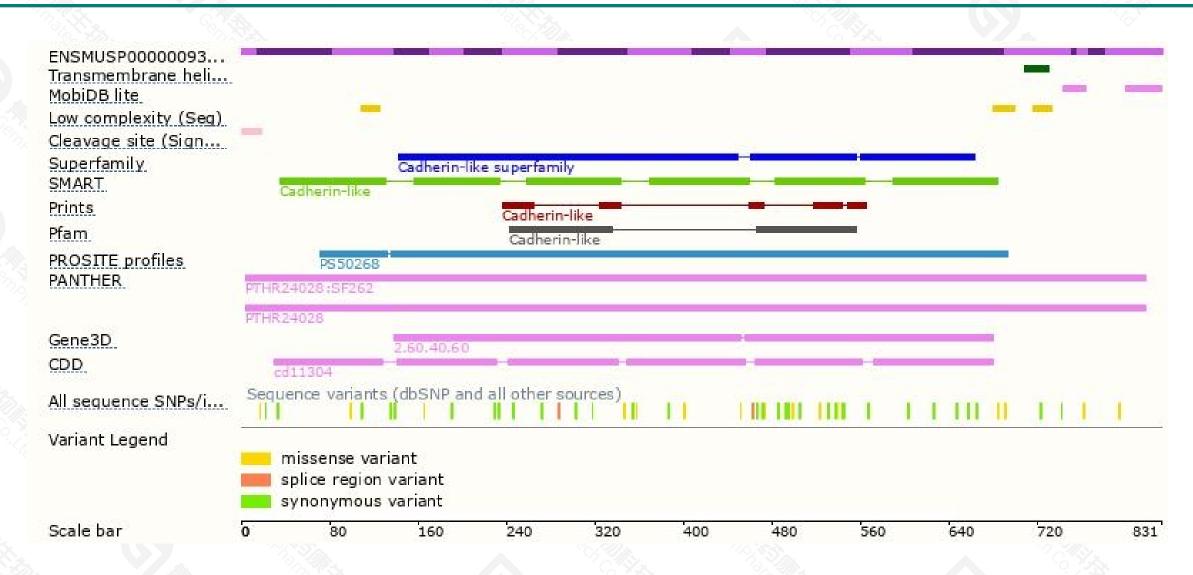
Genomic location distribution





Protein domain







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

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