

Dhrs7c Cas9-KO Strategy

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Project Overview



Project Name Dhrs7c

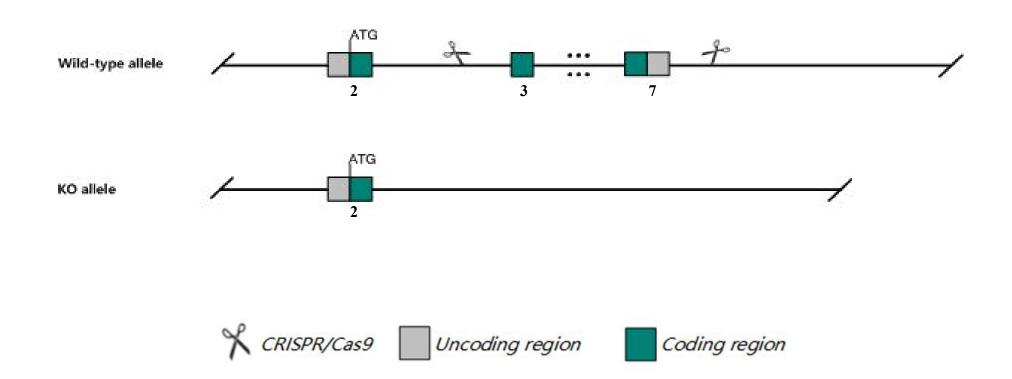
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Dhrs7c* gene. The schematic diagram is as follows:



Technical routes



The *Dhrs7c* gene has 5 transcripts. According to the structure of *Dhrs7c* gene, exon3-exon7 of *Dhrs7c-205* (ENSMUST00000168612.7) transcript is recommended as the knockout region. The region contains 782bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Dhrs7c* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



The *Dhrs7c* gene is located on the Chr11. 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



Dhrs7c dehydrogenase/reductase (SDR family) member 7C [Mus musculus (house mouse)]

Gene ID: 68460, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Dhrs7c provided by MGI

Official Full Name dehydrogenase/reductase (SDR family) member 7C provided by MGI

Primary source MGI:MGI:1915710

See related Ensembl:ENSMUSG00000033044

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 1110001P11Rik, Al120487, SRP-35

Expression Biased expression in heart adult (RPKM 73.2) and mammary gland adult (RPKM 9.8)See more

Orthologs <u>human all</u>

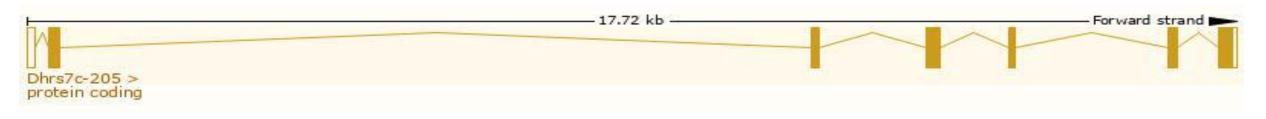
Transcript information Ensembl



The gene has 5 transcripts, all transcripts are shown below:

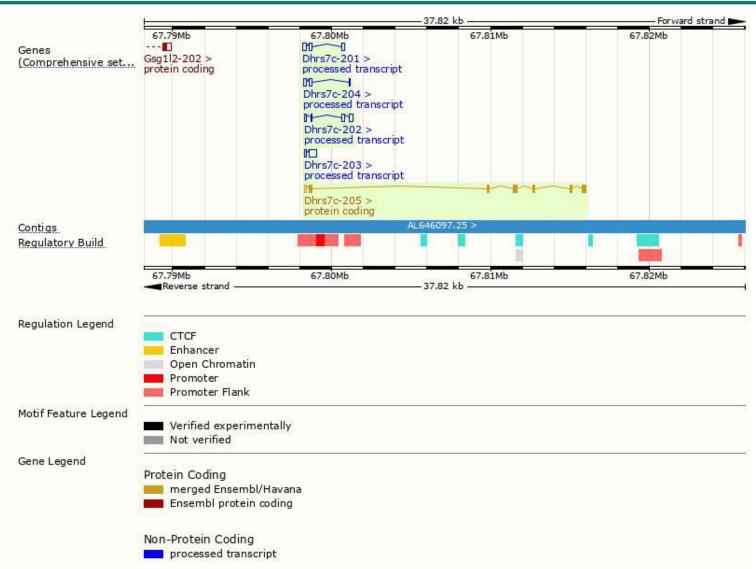
Name	Transcript ID	bp	Protein	Biotype	ccds	UniProt	Flags
Dhrs7c-205	ENSMUST00000168612.7	1108	<u>311aa</u>	Protein coding	CCDS48822	Q8CHS7	TSL:5 GENCODE basic APPRIS P1
Dhrs7c-202	ENSMUST00000135830.1	585	No protein	Processed transcript	-	13 7	TSL:1
Dhrs7c-203	ENSMUST00000142686.1	574	No protein	Processed transcript		94	TSL:2
Dhrs7c-201	ENSMUST00000123862.7	549	No protein	Processed transcript	2	02	TSL:3
Dhrs7c-204	ENSMUST00000149722.5	342	No protein	Processed transcript	5	15	TSL:2

The strategy is based on the design of *Dhrs7c-205* transcript, The transcription is shown below



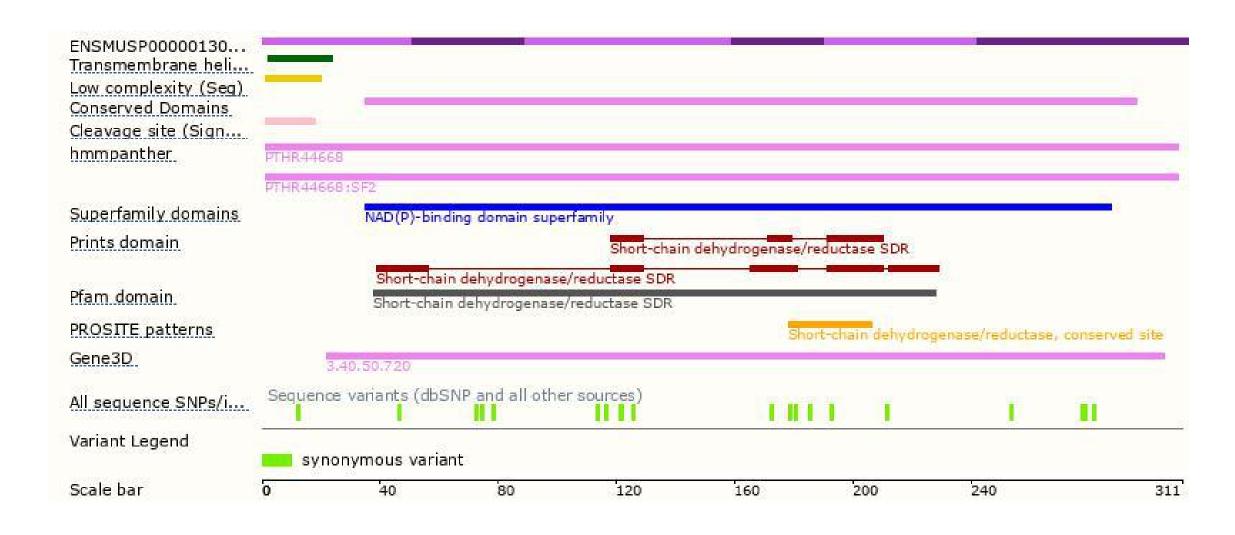
Genomic location distribution





Protein domain







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





