

Dchs2 Cas9-CKO Strategy

Designer:Shuang Zhang

Reviewer: Yun Li

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Project Overview



Project Name Dchs2

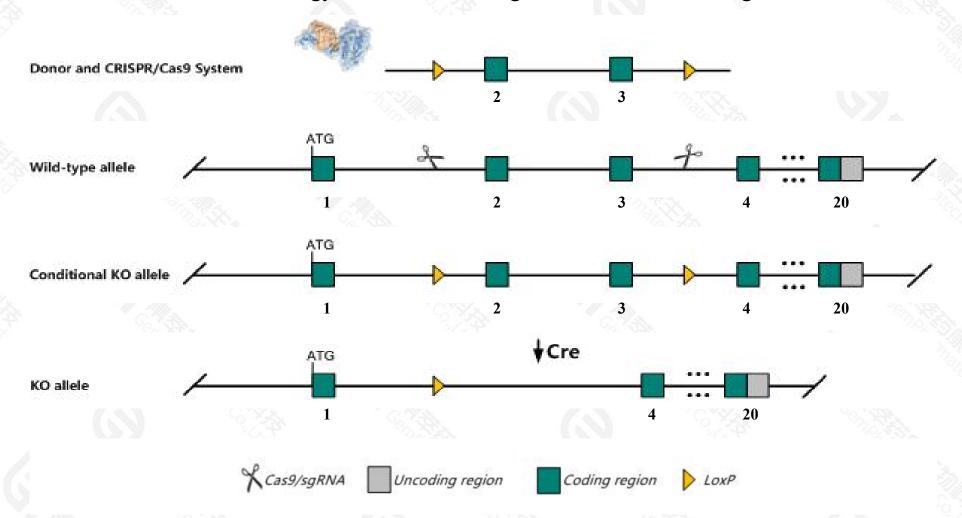
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Dchs2* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Dchs2* gene has 1 transcript. According to the structure of *Dchs2* gene, exon2-exon3 of *Dchs2*201(ENSMUST00000191829.2) transcript is recommended as the knockout region. The region contains 424bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Dchs2* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor was constructed.Cas9, sgRNA 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 was 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



- > The *Dchs2* 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)



Dchs2 dachsous cadherin related 2 [Mus musculus (house mouse)]

Gene ID: 100534287, updated on 25-Sep-2020

Summary

☆ ?

Official Symbol Dchs2 provided by MGI

Official Full Name dachsous cadherin related 2 provided by MGI

Primary source MGI:MGI:2685263

See related Ensembl:ENSMUSG00000102692

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 EG627094, Gm417, Gm6731

Orthologs <u>human</u> all

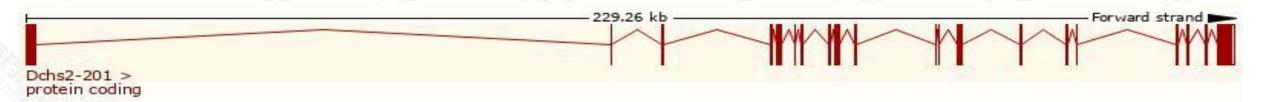
Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

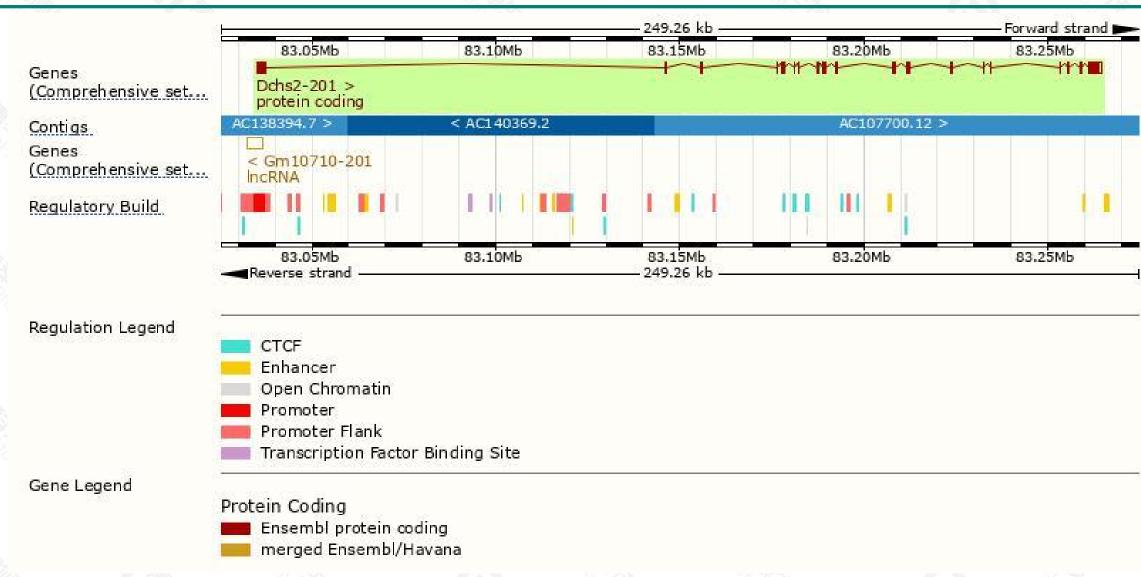
٥	Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
	Dchs2-201	ENSMUST00000191829.2	10783	3346aa	Protein coding	-		TSL:5 , GENCODE basic , APPRIS P1 ,

The strategy is based on the design of *Dchs2-201* transcript, the transcription is shown below:



Genomic location distribution





Protein domain



ENSMUSP00000141... Transmembrane heli... MobiDB lite Low complexity (Seq) Coiled-coils (Ncoils) Superfamily Cadherin-like superfamily SMART Prints Pfam PROSITE profiles Cadherin-like PROSITE patterns PANTHER PTHR24025 PTHR24025:SF20 Gene3D 2,60,40,60 CDD cd11304 Sequence variants (dbSNP and all other sources) All sequence SNPs/i... Variant Legend stop gained inframe deletion missense variant synonymous variant Scale bar 400 800 2000 2400 3346 1200 1600 2800



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

Tel: 025-5864 1534





