

Igdcc4 Cas9-CKO Strategy

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Project Overview



Project Name

Igdcc4

Project type

Cas9-CKO

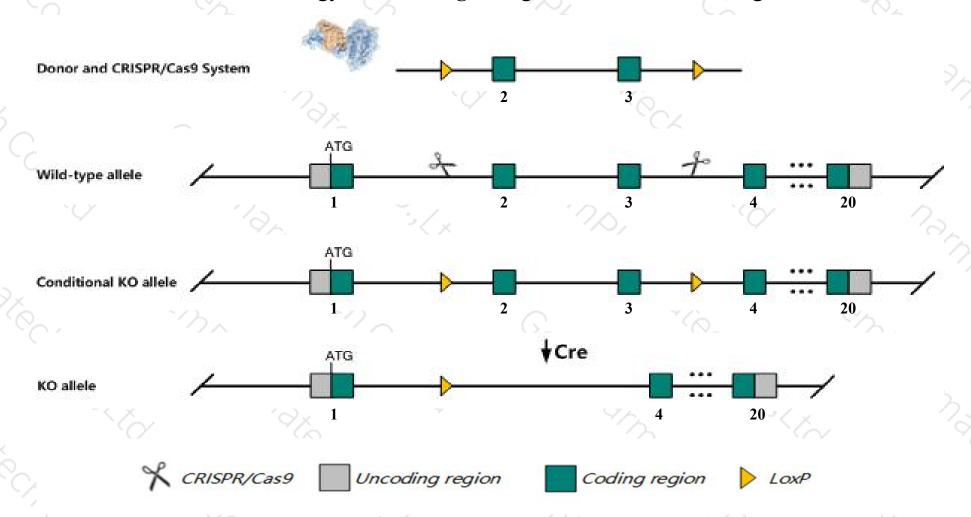
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Igdcc4* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Igdcc4* gene has 7 transcripts. According to the structure of *Igdcc4* gene, exon2-exon3 of *Igdcc4-201*(ENSMUST00000035499.4) transcript is recommended as the knockout region. The region contains 496bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Igdcc4* 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



- ➤ The *Igdcc4* gene is located on the Chr9. 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)



Igdcc4 immunoglobulin superfamily, DCC subclass, member 4 [Mus musculus (house mouse)]

Gene ID: 56741, updated on 24-Oct-2019

Summary

Official Symbol Igdcc4 provided by MGI

Official Full Name immunoglobulin superfamily, DCC subclass, member 4 provided by MGI

Primary source MGI:MGI:1858497

See related Ensembl: ENSMUSG00000032816

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 Nope: DDM36

Expression Broad expression in limb E14.5 (RPKM 23.8), ovary adult (RPKM 16.8) and 21 other tissues See more

Orthologs human all

Genomic context

See Igdcc4 in Genome Data Viewe

Location: 9:9 C

Exon count: 23

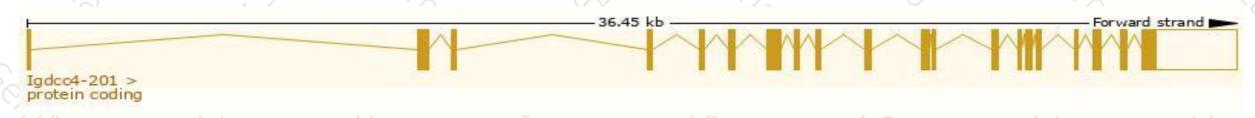
Transcript information (Ensembl)



The gene has 7 transcripts, all transcripts are shown below:

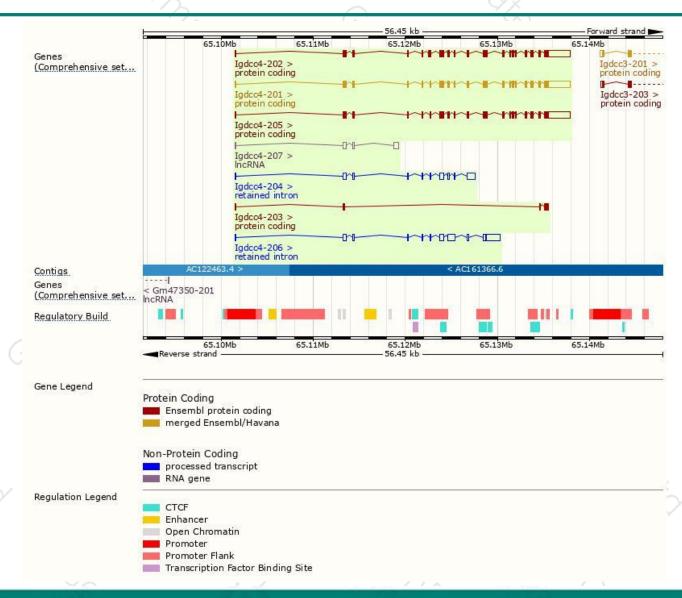
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
lgdcc4-201	ENSMUST00000035499.4	6223	<u>1253aa</u>	Protein coding	CCDS72258	Q9EQS9	TSL:1 GENCODE basic APPRIS P2
lgdcc4-202	ENSMUST00000077696.12	6361	<u>1299aa</u>	Protein coding	8-	E9QAQ0	TSL:5 GENCODE basic
lgdcc4-205	ENSMUST00000213533.1	6220	<u>1252aa</u>	Protein coding	<u>-</u>	Q9EQS9	TSL:1 GENCODE basic APPRIS ALT2
lgdcc4-203	ENSMUST00000166273.1	778	241aa	Protein coding	12	A0JNS2	TSL:1 GENCODE basic
lgdcc4-206	ENSMUST00000214978.1	4130	No protein	Retained intron	7	56	TSL:2
lgdcc4-204	ENSMUST00000213423.1	2572	No protein	Retained intron	2-	-8	TSL:5
lgdcc4-207	ENSMUST00000216542.1	1131	No protein	IncRNA	-	20	TSL:1

The strategy is based on the design of *Igdcc4-201* transcript, The transcription is shown below



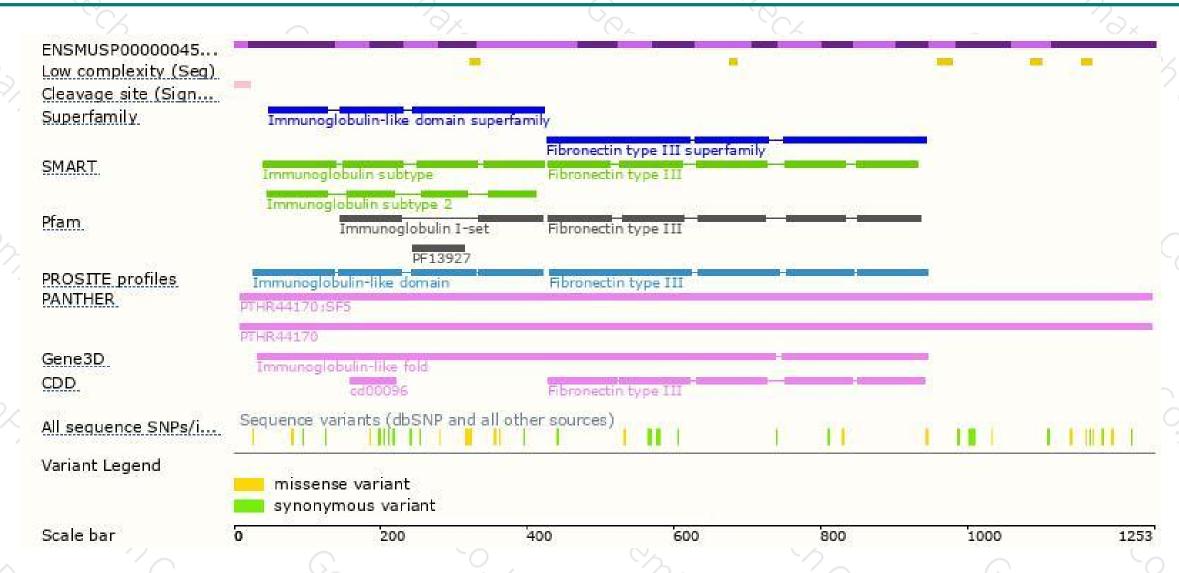
Genomic location distribution





Protein domain







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





