

Isl2 Cas9-CKO Strategy

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Project Overview



Project Name Isl2

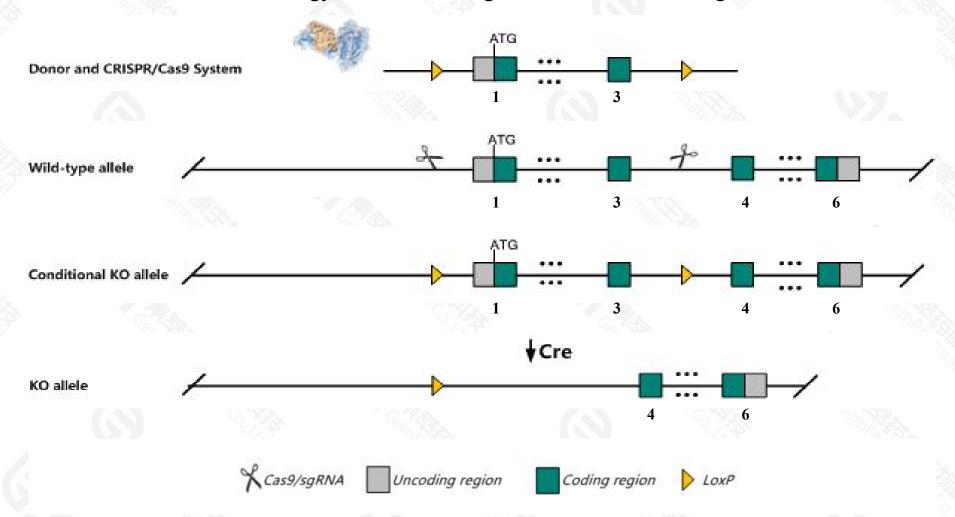
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Isl2* gene. The schematic diagram is as follows:



Technical routes



The *Isl2* gene has 5 transcripts. According to the structure of *Isl2* gene, exon1-exon3 of *Isl2-*201(ENSMUST00000034869.11) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Isl2* 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



According to the existing MGI data, mutations of this gene result in neonatal lethality, motor neuron migration defects and impaired visceral motor neuron differentiation.

The *Isl2* 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



Isl2 insulin related protein 2 (islet 2) [Mus musculus (house mouse)]

Gene ID: 104360, updated on 12-Jan-2021

Summary

☆ ?

Official Symbol Isl2 provided by MGI

Official Full Name insulin related protein 2 (islet 2) provided by MGI

Primary source MGI:MGI:109156

See related Ensembl: ENSMUSG00000032318

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 3110001N10Rik, isle

Expression Biased expression in CNS E11.5 (RPKM 2.3), colon adult (RPKM 1.4) and 11 other tissuesSee more

Orthologs <u>human all</u>

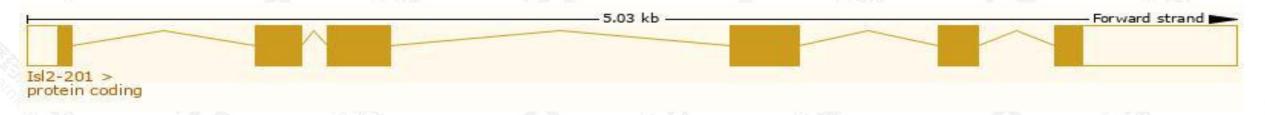
Transcript information Ensembl



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

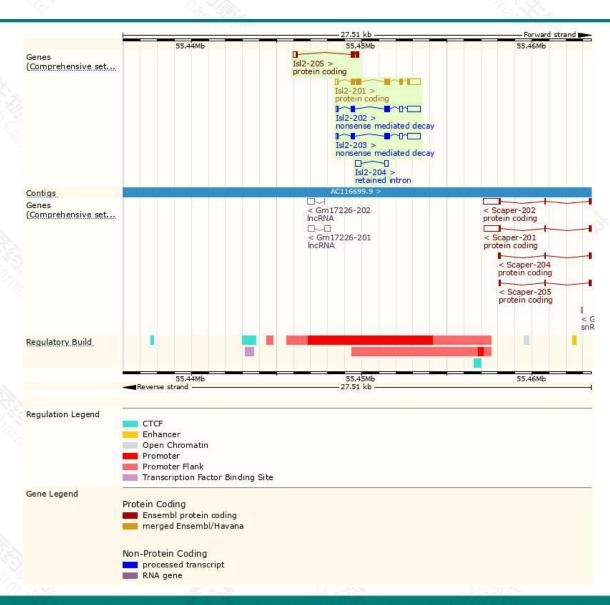
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Isl2-201	ENSMUST00000034869.11	1854	359aa	Protein coding	CCDS23205		TSL:1 , GENCODE basic , APPRIS P1
Isl2-205	ENSMUST00000175950.8	637	<u>151aa</u>	Protein coding	121		CDS 3' incomplete , TSL:3 ,
Isl2-203	ENSMUST00000164373.8	1535	<u>174aa</u>	Nonsense mediated decay	188		TSL:1,
Isl2-202	ENSMUST00000114290.4	1532	<u>174aa</u>	Nonsense mediated decay	6.70		TSL:1,
Isl2-204	ENSMUST00000165302.2	529	No protein	Retained intron	848		TSL:3,

The strategy is based on the design of *Isl2-201* transcript, the transcription is shown below:



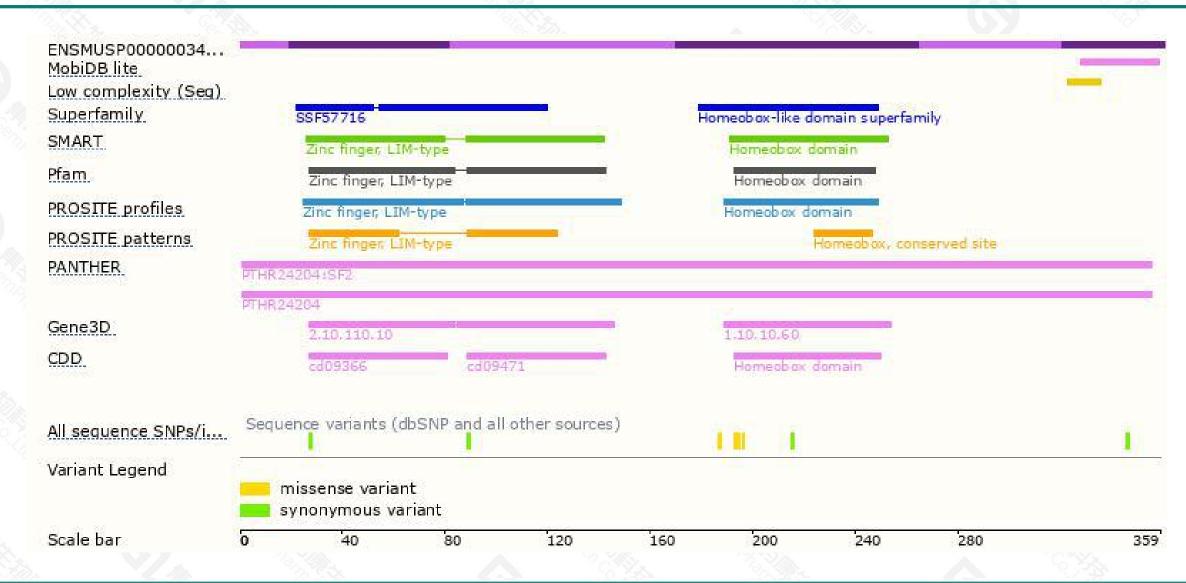
Genomic location distribution





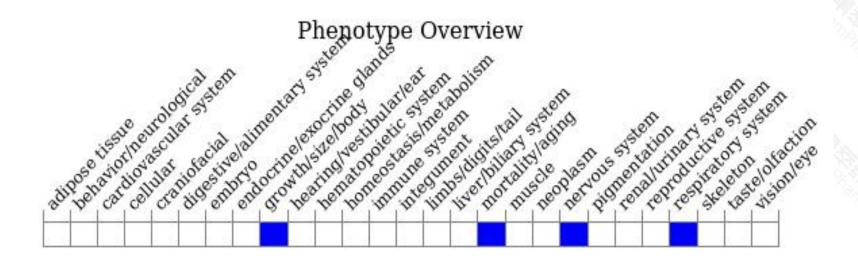
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

According to the existing MGI data, mutations of this gene result in neonatal lethality, motor neuron migration defects and impaired visceral motor neuron differentiation.



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

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