

Tcf7l2 Cas9-KO Strategy

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Project Overview



Project Name Tcf712

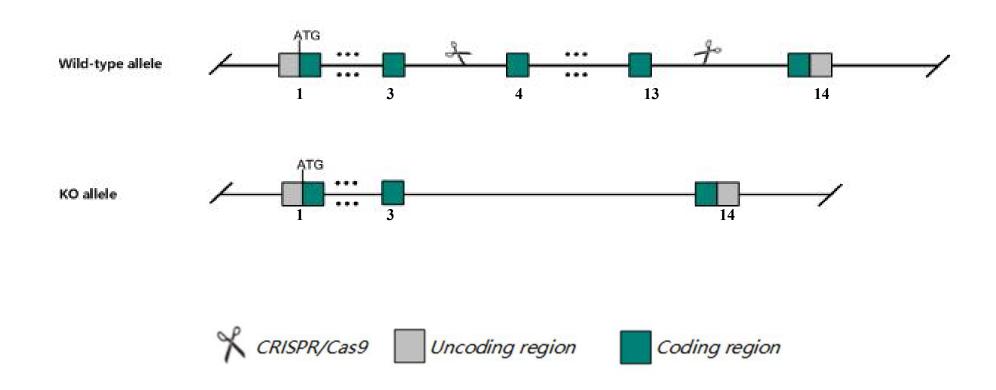
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Tcf7l2* gene. The schematic diagram is as follows:



Technical routes



The *Tcf7l2* gene has 22 transcripts. According to the structure of *Tcf7l2* gene, exon4-exon13 of *Tcf7l2-209* (ENSMUST00000111656.7) transcript is recommended as the knockout region. The region contains 992bp coding sequence.

Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify Tcf712 gene. The brief process is as follows: CRISPR/Cas9 system

Notice



According to the existing MGI data, Animals homozygous for a targeted mutation exhibit intestinal epithelia abnormalities and die shortly after birth. Mice heterozygous for some mutations display abnormalities in glucose homeostasis.

The *Tcf7l2* gene is located on the Chr19. 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



Tcf7l2 transcription factor 7 like 2, T cell specific, HMG box [Mus musculus (house mouse)]

Gene ID: 21416, updated on 5-Mar-2019

Summary

☆ ?

Official Symbol Tcf7I2 provided by MGI

Official Full Name transcription factor 7 like 2, T cell specific, HMG box provided by MGI

Primary source MGI:MGI:1202879

See related Ensembl:ENSMUSG00000024985

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 TCF4B, TCF4E, Tcf-4, Tcf4

Expression Broad expression in whole brain E14.5 (RPKM 35.3), CNS E14 (RPKM 21.4) and 25 other tissuesSee more

Orthologs <u>human</u> all

Transcript information Ensembl



The gene has 22 transcripts, all transcripts are shown below:

The strategy is based on the design of *Tcf712-209* transcript, The transcription is shown below

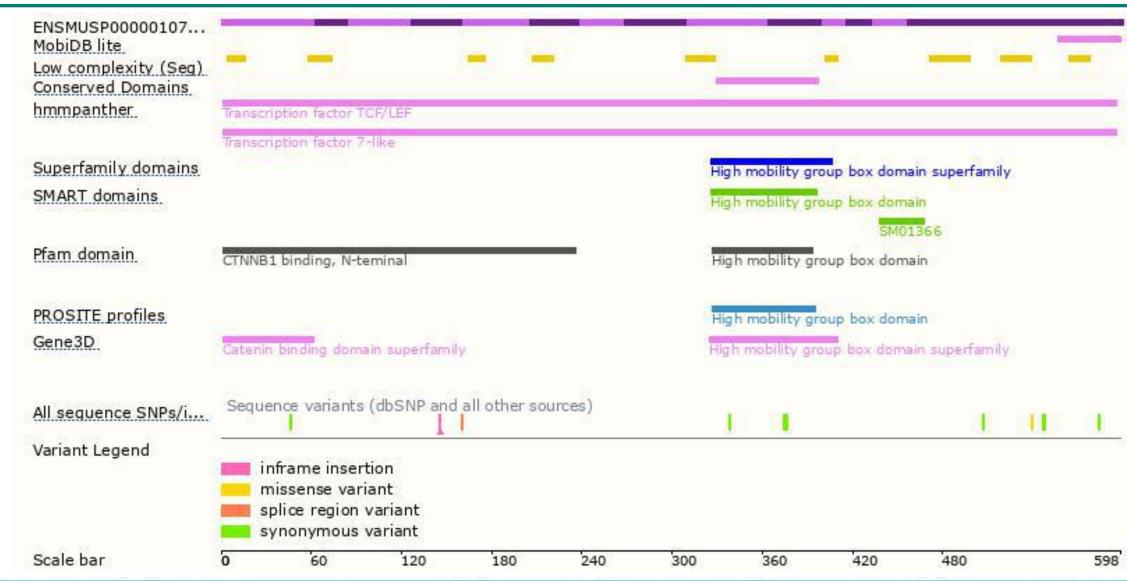


Genomic location distribution



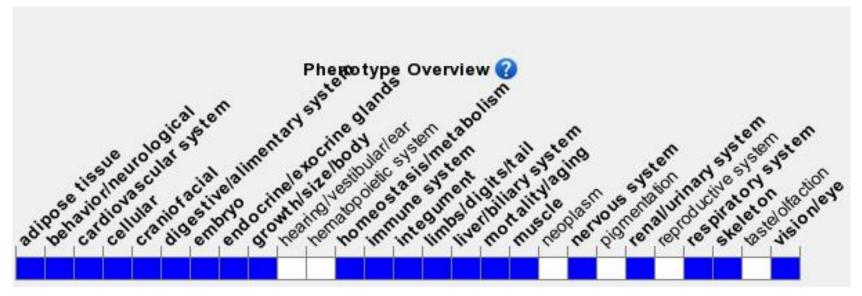
Protein domain





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, Animals homozygous for a targeted mutation exhibit intestinal epithelia abnormalities and die shortly after birth. Mice heterozygous for some mutations display abnormalities in glucose homeostasis



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

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