

Rfx6 Cas9-KO Strategy

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Project Overview



Project Name Rfx6

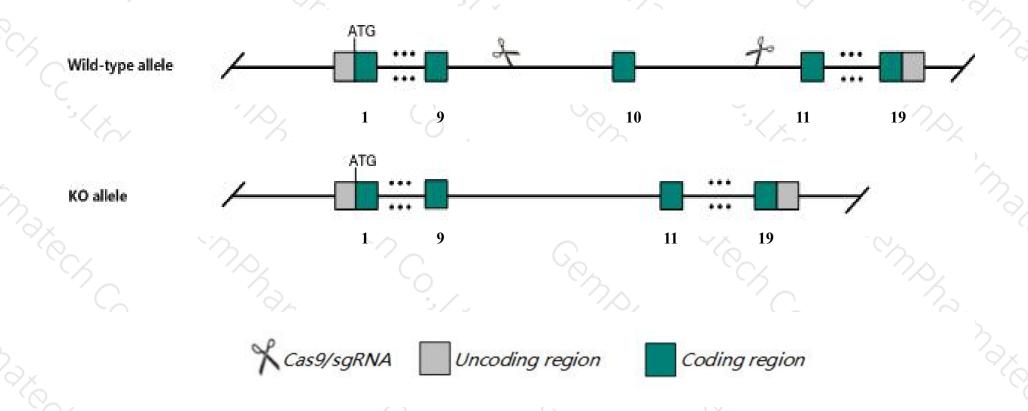
Project type Cas9-KO

Strain background C57BL/6J

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Rfx6* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Rfx6* gene has 5 transcripts. According to the structure of *Rfx6* gene, exon10 of *Rfx6-202* (
 ENSMUST00000122922.9) transcript is recommended as the knockout region. The region contains 50bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Rfx6* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

Notice



- ➤ According to the existing MGI data, Homozygotes fail to feed normally, show small bowel obstruction and die within 2 days of birth. Mutants fail to generate any of the normal islet cell types except for pancreatic-polypeptide-producing cells. Some display a reduced pancreas size; however, primary cilia formation in islets is normal.
- ➤ The *Rfx6* gene is located on the Chr10. 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 gene transcription and translation processes, all risks cannot be predicted under existing information.

Gene information (NCBI)



Rfx6 regulatory factor X, 6 [Mus musculus (house mouse)]

Gene ID: 320995, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Rfx6 provided by MGI

Official Full Name regulatory factor X, 6 provided by MGI

Primary source MGI:MGI:2445208

See related Ensembl:ENSMUSG00000019900

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 4930572007Rik, Rfxdc1

Expression Low expression observed in reference datasetSee more

Orthologs <u>human</u> all

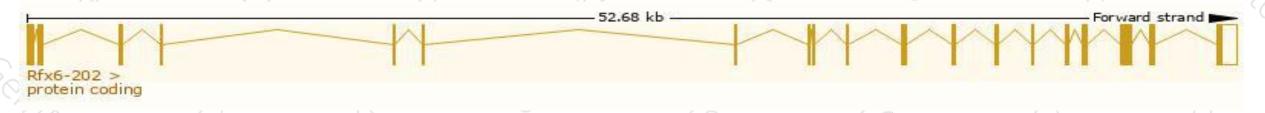
Transcript information (Ensembl)



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

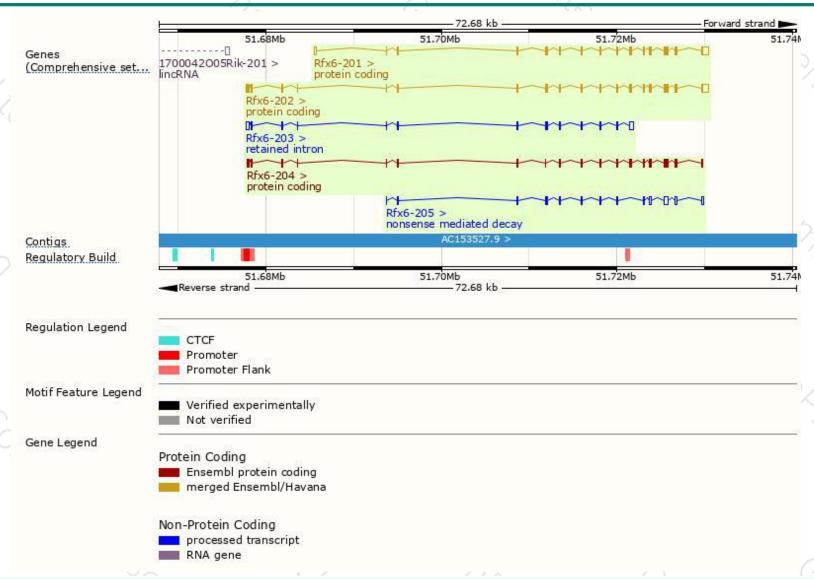
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rfx6-202	ENSMUST00000122922.9	3464	927aa	Protein coding	CCDS48558	Q8C7R7	TSL:5 GENCODE basic APPRIS P1
Rfx6-201	ENSMUST00000050455.4	3088	<u>663aa</u>	Protein coding	CCDS23835	Q8C7R7	TSL:1 GENCODE basic
Rfx6-204	ENSMUST00000219364.1	2682	893aa	Protein coding	2	D5FLC2	TSL:1 GENCODE basic
Rfx6-205	ENSMUST00000219771.1	2101	270aa	Nonsense mediated decay	24	A0A1W2P7F2	CDS 5' incomplete TSL:5
Rfx6-203	ENSMUST00000217662.1	1896	No protein	Retained intron	-	(5)	TSL:1

The strategy is based on the design of Rfx6-202 transcript, The transcription is shown below



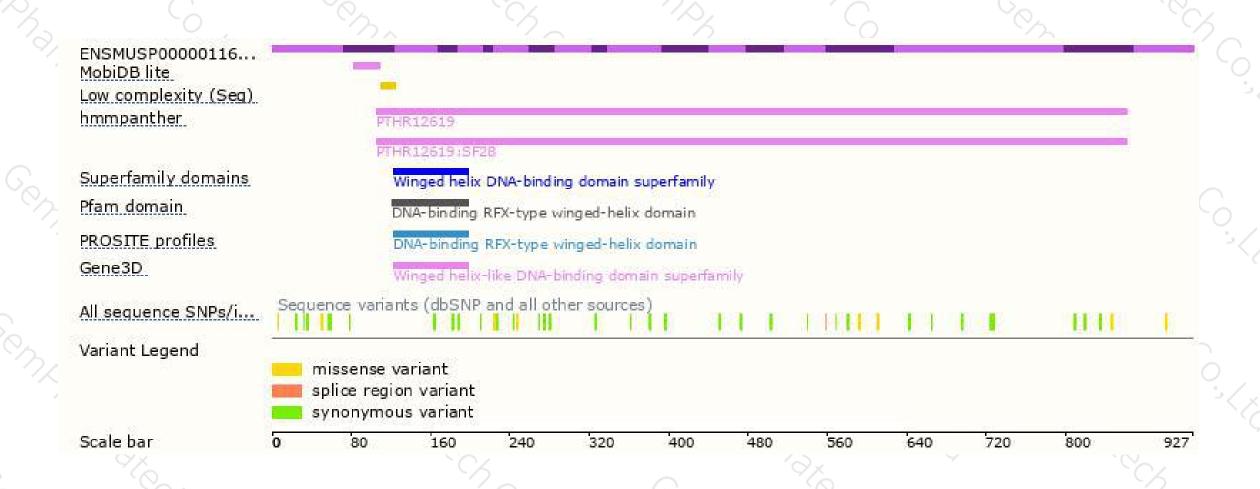
Genomic location distribution





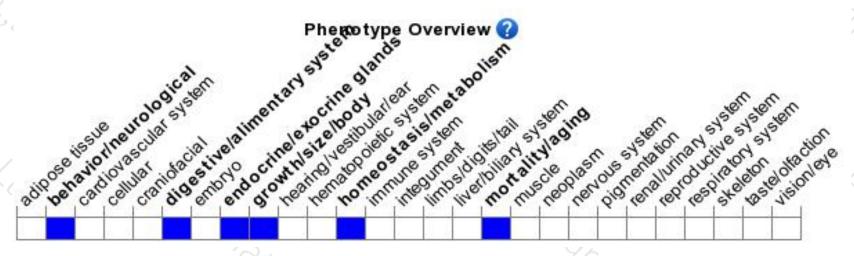
Protein domain





Mouse phenotype description(MGI)





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

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

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