Htr1b Cas9-KO Strategy

Designer: Huan Wang

Design Date: 2020-7-22

Project Overview



Project Name

Htr1b

Project type

Cas9-KO

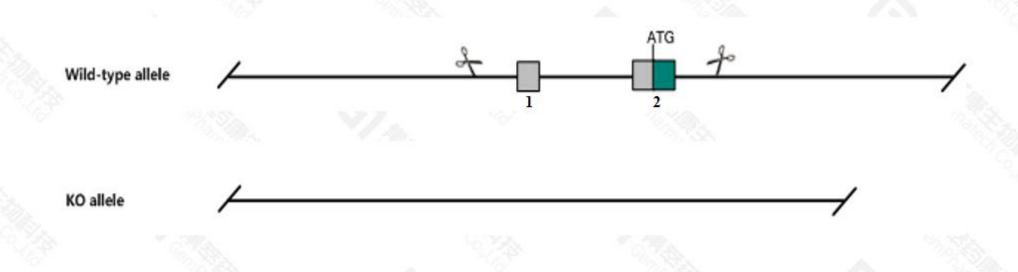
Strain background

C57BL/6J

Conditional Knock Out strategy



This model will use CRISPR-Cas9 technology to edit the *Htr1b* gene. The schematic diagram is as follows:



CRISPR-Cas9 Noncoding region Coding region

Technical routes



The *Htr1b* gene has 2 transcript. According to the structure of *Htr1b* gene, exon 1-2 of *Htr1b*-202 transcript is recommended as the knock Out region. The region contains all coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify xxx 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, mice homozygous for a knock-out allele exhibit an increase in body weight, aggression, drinking behavior, and osteoblast proliferation and bone mass, and show altered spatial learning and operant conditional behavior as well as reduced anxiety-related response and startle reflex, and small testes.
- The KO region contains functional region of the *D430036J16RIK* gene. Knock Out the region may affect the function of *D430036J16RIK* gene.
- The Htr1b gene is located on the Chr9. If the knock out 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Htr1b 5-hydroxytryptamine (serotonin) receptor 1B [Mus musculus (house mouse)]

Gene ID: 15551, updated on 12-Feb-2019

Summary

☆ ?

Official Symbol Htr1b provided by MGI

Official Full Name 5-hydroxytryptamine (serotonin) receptor 1B provided by MGI

Primary source MGI:MGI:96274

See related Ensembl: ENSMUSG00000049511

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae;

Mus; Mus

Also known as 5-HT-1B

Orthologs human all

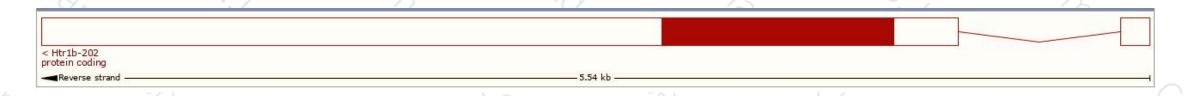
Transcript information (Ensembl)



The gene has 2 transcripts, and all transcripts are shown below:

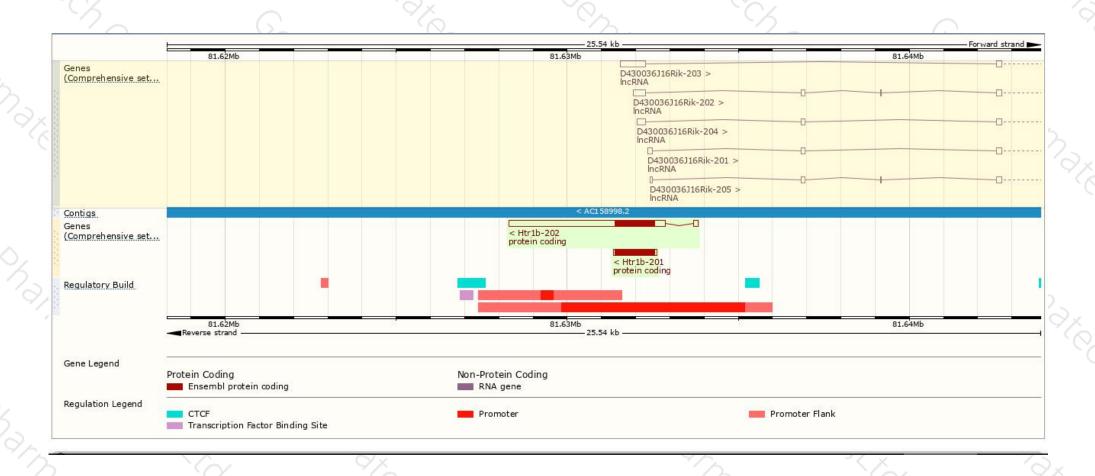
Name 🍦	Transcript ID v	bp 🌲	Protein 4	Biotype 🍦	CCDS .	UniProt	Flags		
Htr1b-202	ENSMUST00000183482.1	4724	386aa	Protein coding	CCDS23370 ₽	P28334₺ Q0VES5₺	TSL:3	GENCODE basic	APPRIS P1
Htr1b-201	ENSMUST00000051005.4	1273	386aa	Protein coding	CCDS23370 ₽	<u>P28334</u>	TSL:NA	GENCODE basic	APPRIS P1

The strategy is based on the design of *Htr1b*-202 transcript, The transcription is shown below



Genomic location distribution

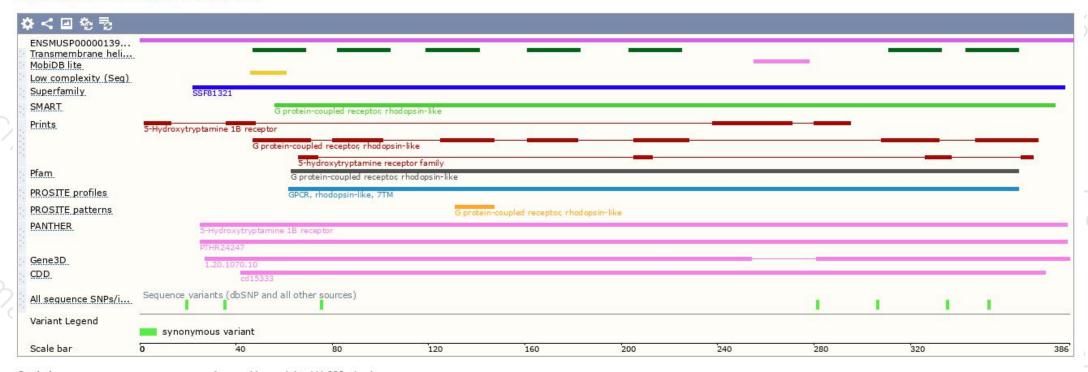




Protein domain



Protein domains for ENSMUSP00000139389.1



Statistics

Ave. residue weight: 111.603 g/mol

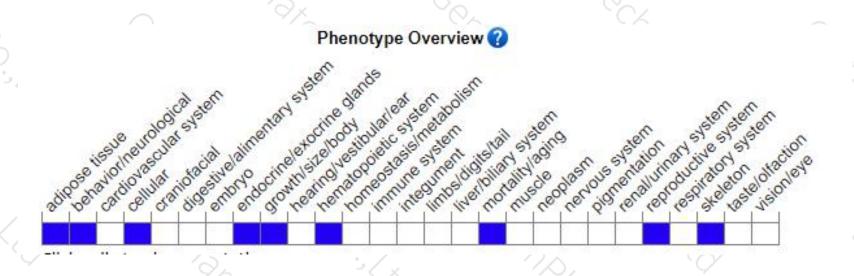
Charge: 12.0

Isoelectric point: 8.7811

Molecular weight: 43,078.68 g/mol Number of residues: 386 aa

Mouse phenotype description(MGI)





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

Mice homozygous for a knock-out allele exhibit an increase in body weight, aggression, drinking behavior, and osteoblast proliferation and bone mass, and show altered spatial learning and operant conditional behavior as well as reduced anxiety-related response and startle reflex, and small testes.

If you have any questions, you are welcome to inquire. Tel: 025-5864 1534





