

Htr5a Cas9-KO Strategy

Designer:

Min Guan

Reviewer:

Yang Zeng

Design Date:

2018-6-28

Project Overview

Project Name

Htr5a

Project type

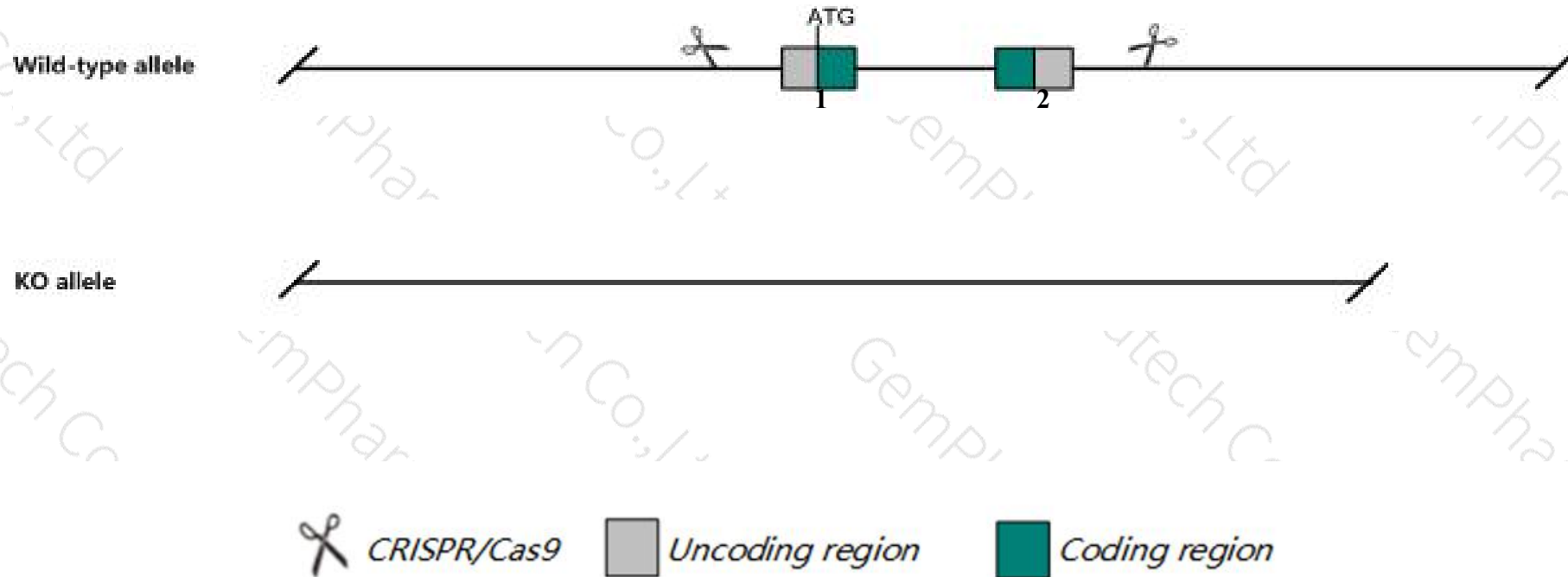
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Htr5a* gene. The schematic diagram is as follows:



- The *Htr5a* gene has 1 transcript. According to the structure of *Htr5a* gene, exon1-exon2 of *Htr5a-201* (ENSMUST00000036227.6) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Htr5a* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, There are conflicting reports about mice with homozygous null alleles regarding increased exploratory behavior and an abnormal exploratory behavior response to LSD.
- The *Htr5a* gene is located on the Chr5. 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)

Htr5a 5-hydroxytryptamine (serotonin) receptor 5A [*Mus musculus* (house mouse)]

Gene ID: 15563, updated on 12-Aug-2019

Summary

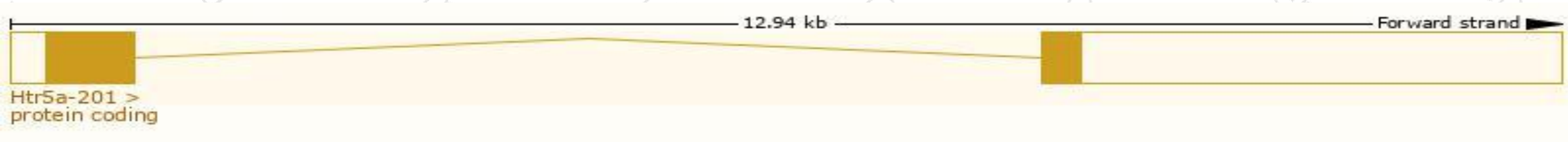
Official Symbol	Htr5a provided by MGI
Official Full Name	5-hydroxytryptamine (serotonin) receptor 5A provided by MGI
Primary source	MGI:MGI:96283
See related	Ensembl:ENSMUSG00000039106
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Htr5
Expression	Biased expression in cortex adult (RPKM 2.1), frontal lobe adult (RPKM 2.0) and 4 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

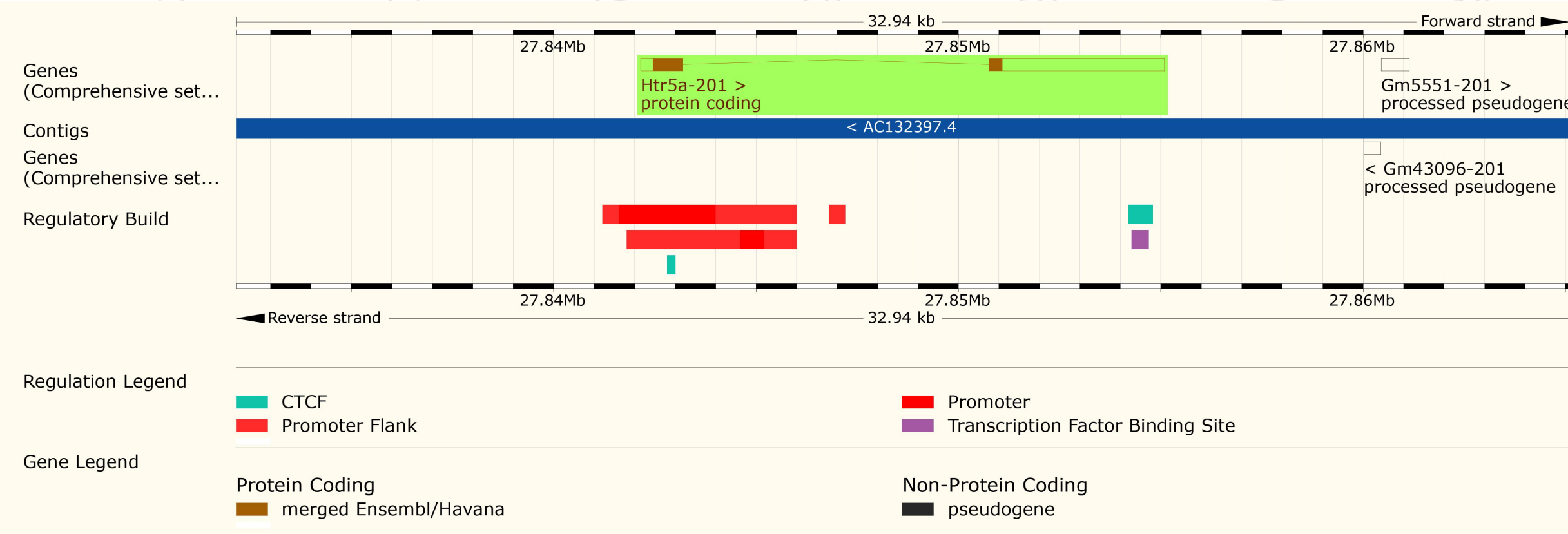
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Htr5a-201	ENSMUST00000036227.6	5376	357aa	Protein coding	CCDS19141	P30966	TSL:1 GENCODE basic APPRIS P1

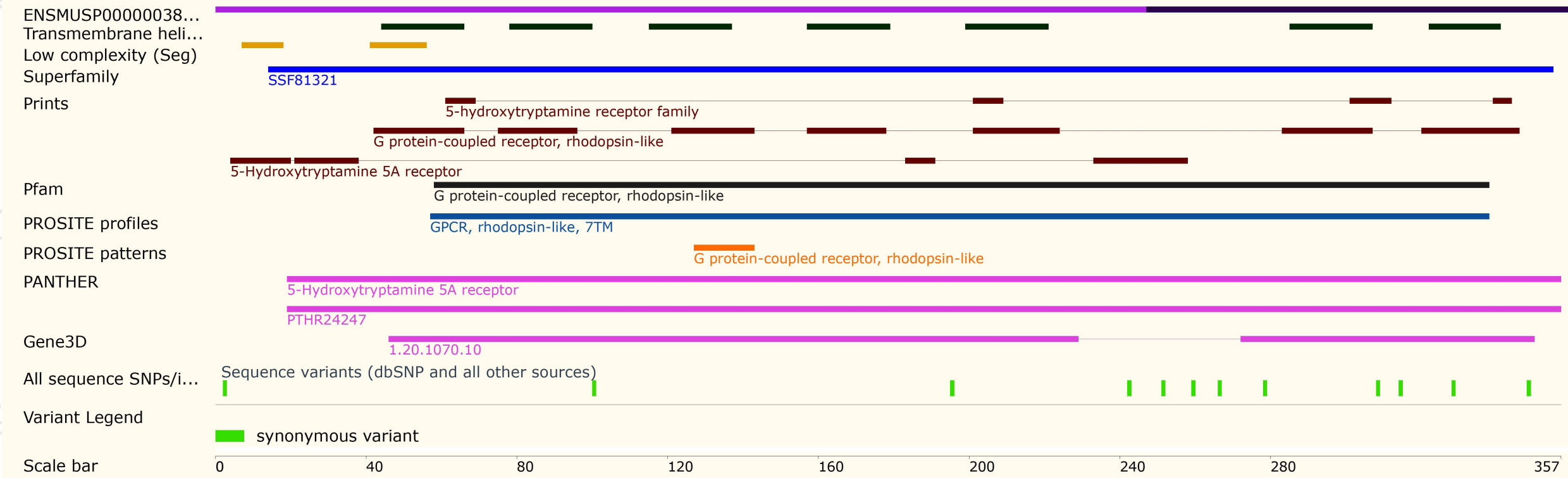
The strategy is based on the design of *Htr5a-201* transcript, The transcription is shown below



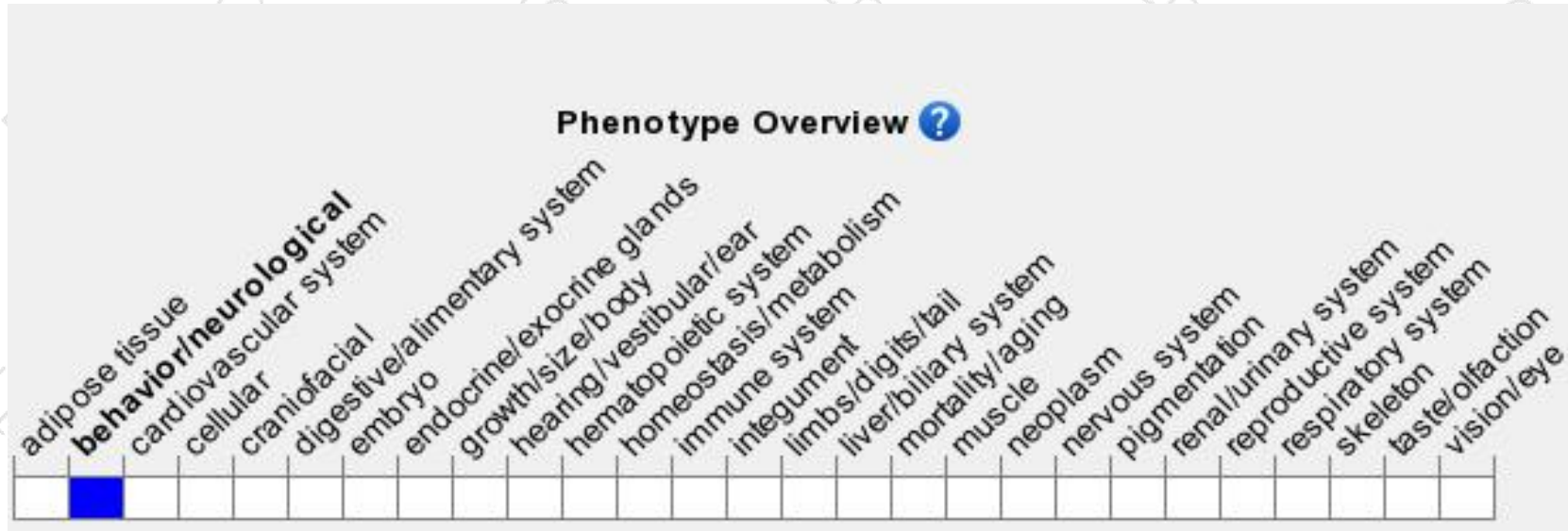
Genomic location distribution



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, There are conflicting reports about mice with homozygous null alleles regarding increased exploratory behavior and an abnormal exploratory behavior response to LSD.

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

Tel: 400-9660890

