

Ntsr1 Cas9-KO Strategy

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Design Date:

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Project Overview



Project Name

Ntsr1

Project type

Cas9-KO

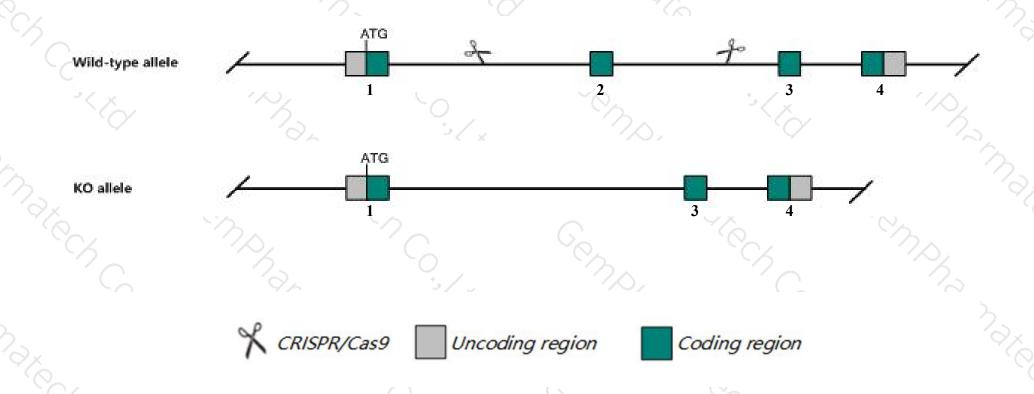
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ntsr1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Ntsr1* gene has 2 transcripts. According to the structure of *Ntsr1* gene, exon2 of *Ntsr1-201*(ENSMUST00000029084.8) transcript is recommended as the knockout region. The region contains 217bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Ntsr1* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ According to the existing MGI data, Mice deficient for this marker have normal baseline prepulse inhibition responses and acoustic startle responses. Mice are heavier, eat more, and have lower body temperatures.
- The *Ntsr1* gene is located on the Chr2. 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)



Ntsr1 neurotensin receptor 1 [Mus musculus (house mouse)]

Gene ID: 18216, updated on 10-Aug-2019

Summary

☆ ?

Official Symbol Ntsr1 provided by MGI

Official Full Name neurotensin receptor 1 provided by MGI

Primary source MGI:MGI:97386

See related Ensembl: ENSMUSG00000027568

Gene type protein coding
RefSeq status PROVISIONAL
Organism Mus musculus

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

Murinae; Mus; Mus

Also known as NTR1; NT-1R; NTR-1

Expression Biased expression in CNS E18 (RPKM 4.6), genital fat pad adult (RPKM 1.5) and 10 other tissues See more

Orthologs human all

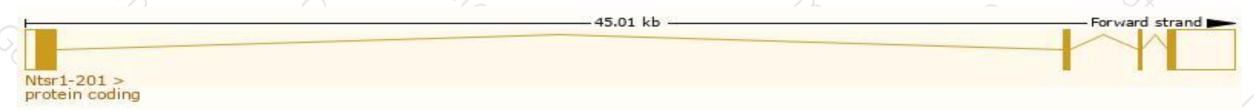
Transcript information (Ensembl)



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

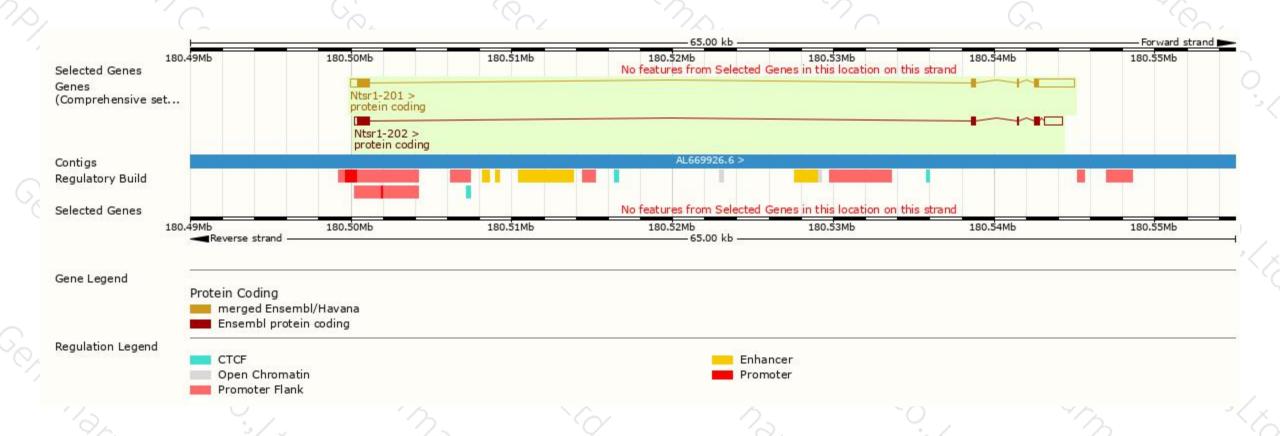
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ntsr1-201	ENSMUST00000029084.8	3917	<u>424aa</u>	Protein coding	CCDS17177	A2ACT4 088319	TSL:1 GENCODE basic APPRIS P1
Ntsr1-202	ENSMUST00000170448.1	2594	<u>424aa</u>	Protein coding	CCDS17177	A2ACT4 088319	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of Ntsr1-201 transcript, The transcription is shown below



Genomic location distribution





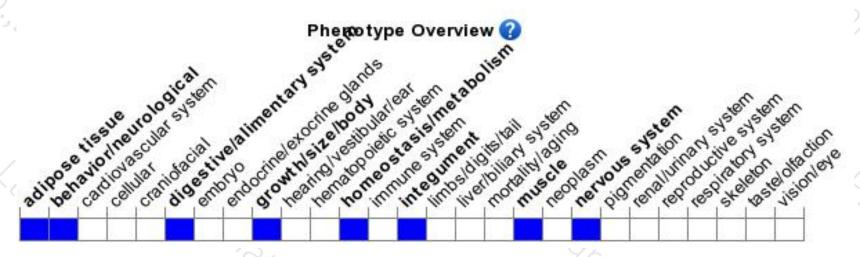
Protein domain





Mouse phenotype description(MGI)





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

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If you have any questions, you are welcome to inquire. Tel: 400-9660890





