

# Nptx2 Cas9-CKO Strategy

**Designer:** 

Yanhua Shen

**Reviewer:** 

**Xueting Zhang** 

**Design Date:** 

2019-12-12

# **Project Overview**



**Project Name** 

Nptx2

**Project type** 

Cas9-CKO

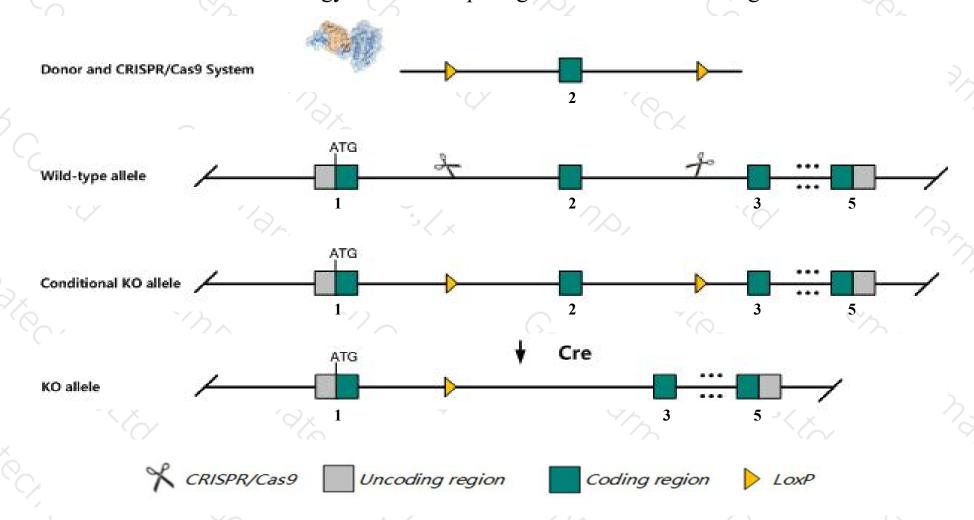
Strain background

C57BL/6JGpt

# Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Nptx2* gene. The schematic diagram is as follows:



### Technical routes



- The *Nptx2* gene has 1 transcript. According to the structure of *Nptx2* gene, exon2 of *Nptx2-201*(ENSMUST00000071782.7) 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 *Nptx2* gene. The brief process is as follows:CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt 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/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

### **Notice**



- > According to the existing MGI data, Mice homozygous for a null mutation of this gene display a mild alteration in retinal ganglion cell innervation but are fertile with no obvious behavioral abnormalities.
- The *Nptx2* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

### Gene information (NCBI)



#### Nptx2 neuronal pentraxin 2 [ Mus musculus (house mouse) ]

Gene ID: 53324, updated on 10-Oct-2019

Summary

△ ?

Official Symbol Nptx2 provided by MGI

Official Full Name neuronal pentraxin 2 provided by MGI

Primary source MGI:MGI:1858209

See related Ensembl: ENSMUSG00000059991

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 np2; Narp

Expression Broad expression in cortex adult (RPKM 14.8), frontal lobe adult (RPKM 13.5) and 17 other tissues See more

Orthologs human all

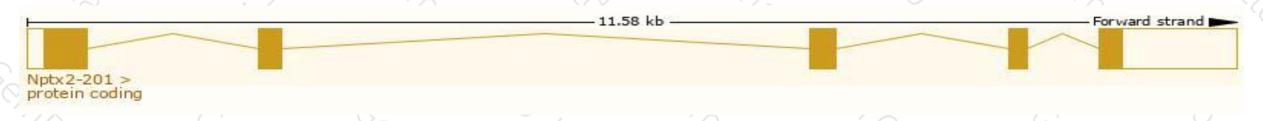
# Transcript information (Ensembl)



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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	l
Nptx2-201	ENSMUST00000071782.7	2536	429aa	Protein coding	CCDS19849	070340	TSL:1 GENCODE basic APPRIS P1	Ľ

The strategy is based on the design of *Nptx2-201* transcript, The transcription is shown below



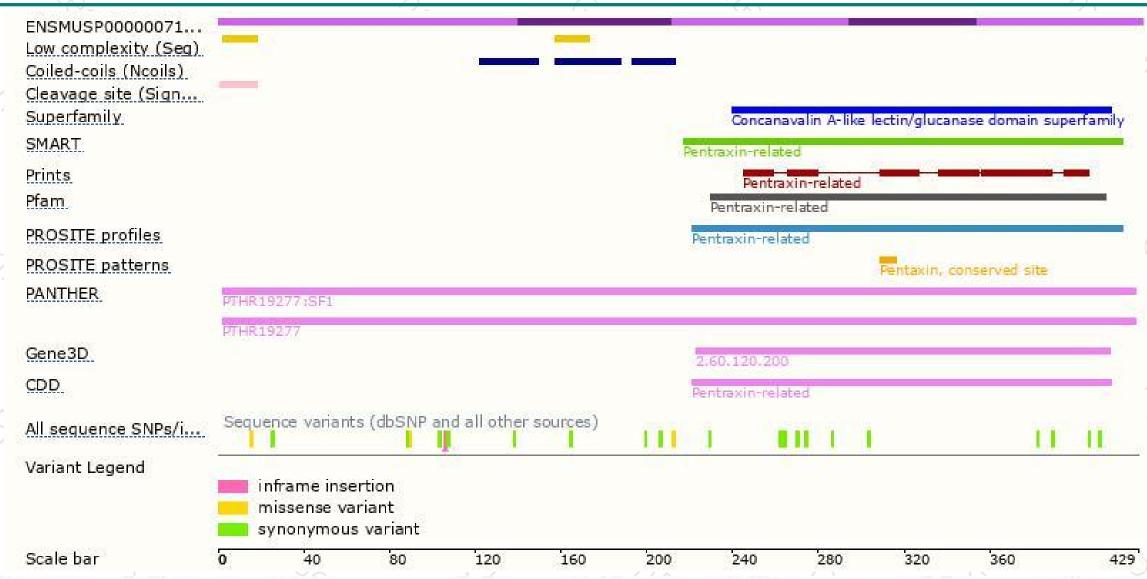
### Genomic location distribution





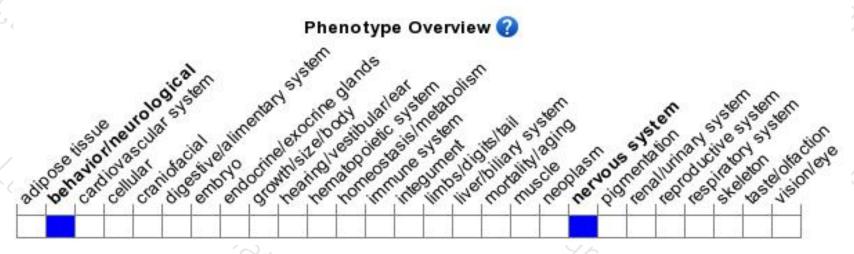
### Protein domain





# 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, Mice homozygous for a null mutation of this gene display a mild alteration in retinal ganglion cell innervation but are fertile with no obvious behavioral abnormalities.



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





