

Ppp2r2c Cas9-KO Strategy

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Project Overview



Project Name

Ppp2r2c

Project type

Cas9-KO

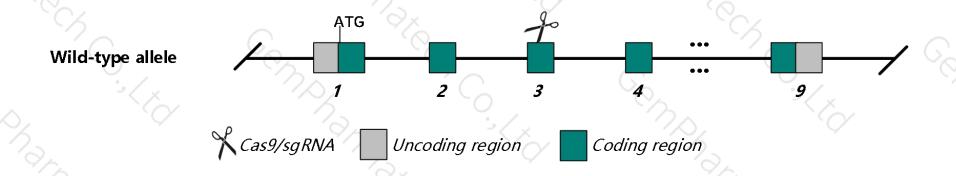
Strain background

C57BL/6N

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ppp2r2c* gene. The schematic diagram is as follows:



Technical routes



➤ In this project we use CRISPR/Cas9 technology to modify *Ppp2r2c* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6N 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/6N mice.

Notice



- ➤ The *Ppp2r2c* 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)



Ppp2r2c protein phosphatase 2, regulatory subunit B, gamma [Mus musculus (house mouse)]

Gene ID: 269643, updated on 12-Aug-2019

Summary

↑ ?

Official Symbol Ppp2r2c provided by MGI

Official Full Name protein phosphatase 2, regulatory subunit B, gamma provided by MGI

Primary source MGI:MGI:2442660

See related Ensembl: ENSMUSG00000029120

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 PR52; IMYPNO; IMYPNO1; 6330548O06Rik

Expression Biased expression in cortex adult (RPKM 105.0), frontal lobe adult (RPKM 88.2) and 6 other tissues See more

Orthologs human all

Transcript information (Ensembl)

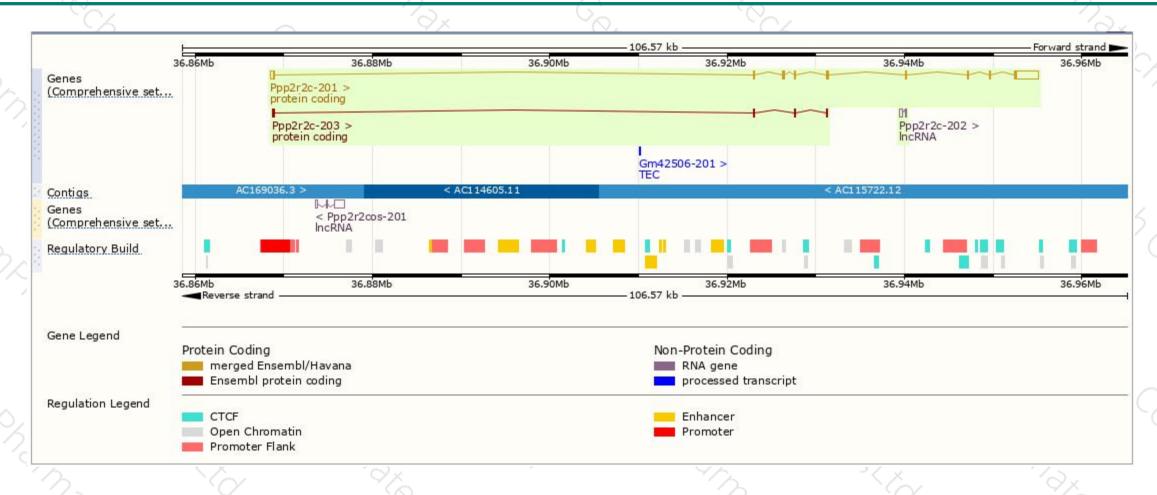


The gene has 3 transcripts, all transcripts are shown below:

Name 🍦	Transcript ID	bp 🌲	Protein 🌲	Biotype 🍦	CCDS 🍦	UniProt 🍦	Flags
Ppp2r2c-201	ENSMUST00000031003.10	4088	447aa	Protein coding	CCDS19244&	Q8BG02₽	TSL:1 GENCODE basic APPRIS P1
Ppp2r2c-203	ENSMUST00000201156.1	461	<u>76aa</u>	Protein coding	-	A0A0J9YUU0₽	TSL:5 GENCODE basic
Ppp2r2c-202	ENSMUST00000138124.1	453	No protein	I IncRNA	-	-	TSL:3

Genomic location distribution







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

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