

Ppp1r1a Cas9-KO Strategy

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Design Date:2019-09-25

Project Overview



Project Name Ppp1r1a

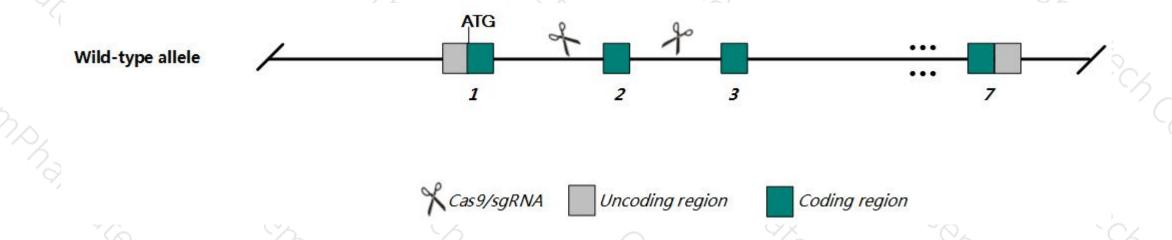
Project type Cas9-KO

Strain background C57BL/6N

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ppp1r1a* gene. The schematic diagram is as follows:



Technical routes



➤ In this project we use CRISPR/Cas9 technology to modify *Ppp1r1a* 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



- ➤ According to MGI, Homozygous mutant mice show decreased long term poteniation, but normal performance in water maze tests.
- The *Ppp1r1a* gene is located on the Chr15, 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)



Ppp1r1a protein phosphatase 1, regulatory inhibitor subunit 1A [Mus musculus (house mouse)]

Gene ID: 58200, updated on 13-Aug-2019

Summary

Official Symbol Ppp1r1a provided by MGI

Official Full Name protein phosphatase 1, regulatory inhibitor subunit 1A provided by MGI

Primary source MGI:MGI:1889595

See related Ensembl: ENSMUSG00000022490

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 I-1: 0610038N18Rik

Expression Biased expression in kidney adult (RPKM 53.9), mammary gland adult (RPKM 24.9) and 11 other tissues See more

Orthologs human all

Transcript information (Ensembl)

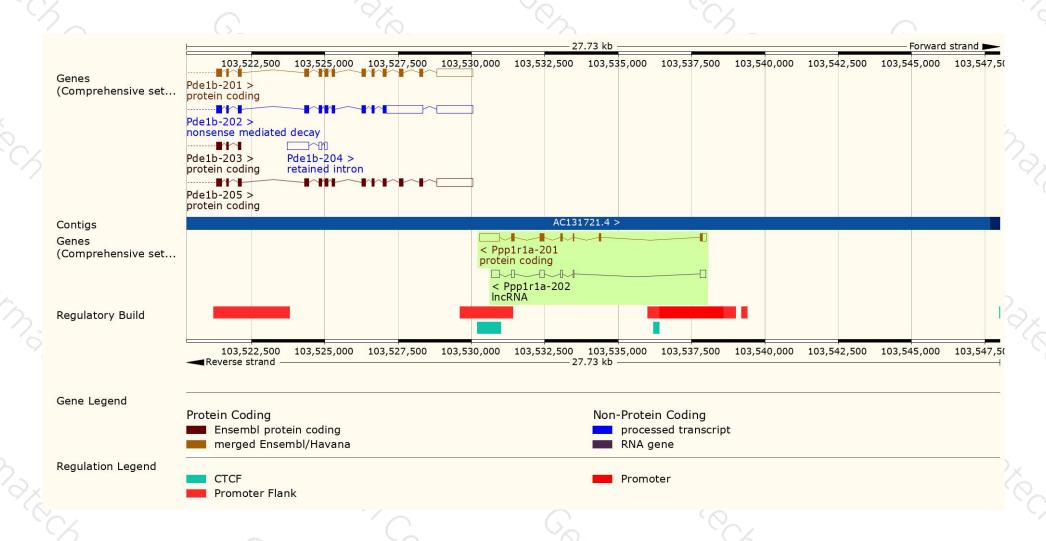


The gene has 2 transcripts, all transcripts are shown below:

Name A	Transcript ID 👙	bp 🌲	Protein 🍦	Biotype 🝦	CCDS 🍦	UniProt 4	Flags		
Ppp1r1a-201	ENSMUST00000023133.7	1306	<u>171aa</u>	Protein coding	CCDS37237 ₺	Q9ERT9₽	TSL:1	GENCODE basic	APPRIS P1
Ppp1r1a-202	ENSMUST00000228813.1	821	No protein	IncRNA	670	5		170	

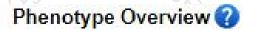
Genomic location distribution

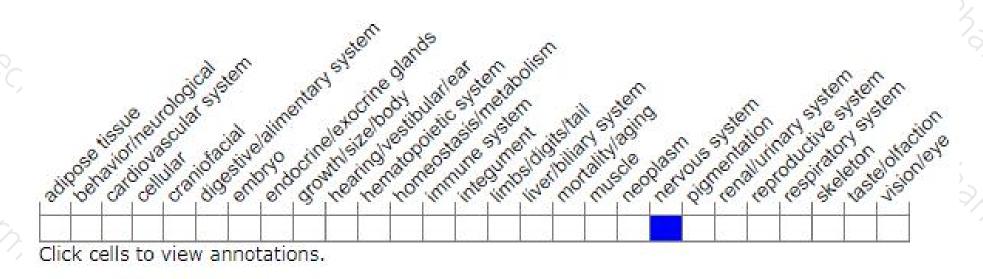




Mouse phenotype description(MGI)







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

Homozygous mutant mice show decreased long term poteniation, but normal performance in water maze tests.



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

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