

Ppp6c Cas9-KO Strategy

Designer: Ruirui Zhang

Reviewer: FengjuanWang

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Project Overview

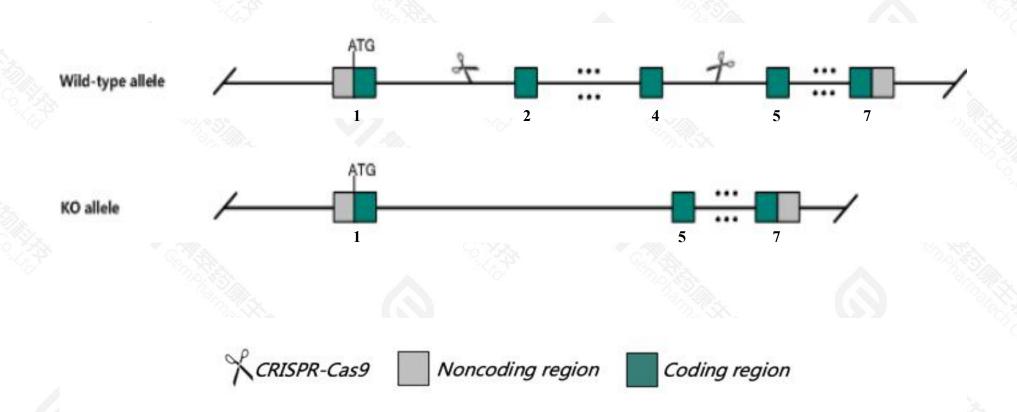


Project Name	Ppp6c
Project type	Cas9-KO
Strain background	C57BL/6JGpt

Knockout strategy



This model will use CRISPR-Cas9 technology to edit the *Ppp6c* gene. The schematic diagram is as follows:



Technical routes



- > The *Ppp6c* gene has 5 transcripts. According to the structure of *Ppp6c* gene, exon2-exon4 of *Ppp6c-201*(ENSMUST00000028087.6) transcript is recommended as the knockout region. The region contains 304bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR-Cas9 technology to modify *Ppp6c* gene. The brief process is as follows: CRISPR-Cas9 system 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.

Notice



- > According to the existing MGI data,mice homozygous for a knock-out allele exhibit abnormal embryonic development and embryonic lethality. Mice homozygous for a conditional allele activated in skin cells exhibit increased susceptibility to chemically induced skin tumors with increased proliferative and inflammatory responses in the skin.
- > The *Ppp6c* 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)



Ppp6c protein phosphatase 6, catalytic subunit [Mus musculus (house mouse)]

Gene ID: 67857, updated on 19-Mar-2019

Summary



Official Symbol Ppp6c provided by MGI

Official Full Name protein phosphatase 6, catalytic subunit provided by MGI

Primary source MGI:MGI:1915107

See related Ensembl: ENSMUSG00000026753

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 2310003C10Rik, Pp6C

Expression Ubiquitous expression in placenta adult (RPKM 11.6), bladder adult (RPKM 11.3) and 28 other tissuesSee more

Orthologs <u>human all</u>

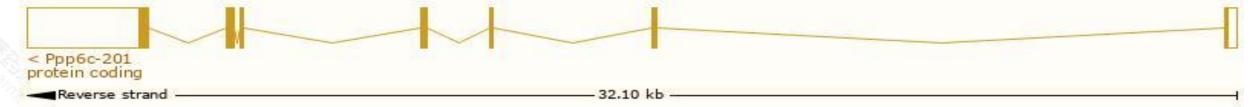
Transcript information (Ensembl)



The gene has 5 transcripts, all transcripts are shown below:

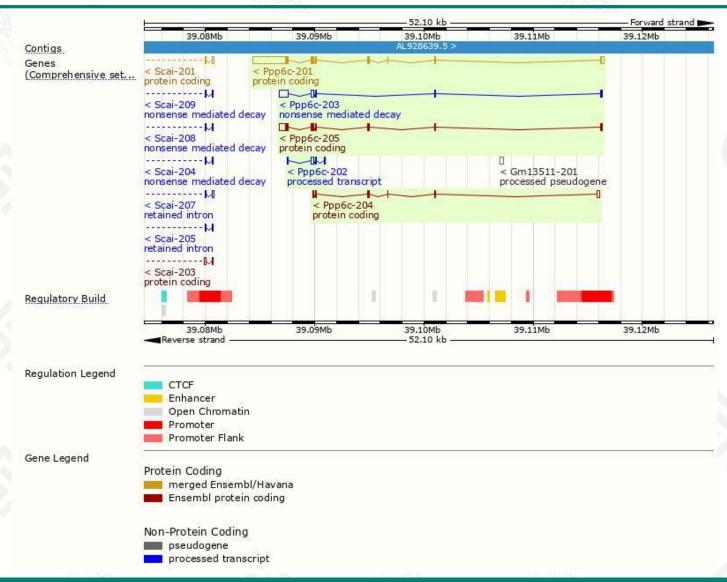
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ppp6c-201	ENSMUST00000028087.5	4115	305aa	Protein coding	CCDS16017	Q9CQR6	TSL:1 GENCODE basic APPRIS P1
Ppp6c-205	ENSMUST00000204701.2	1495	283aa	Protein coding	N 4 3	A0A0N4SVL9	TSL:5 GENCODE basic
Ppp6c-204	ENSMUST00000204368.1	635	<u>83aa</u>	Protein coding	020	A0A0N4SW66	CDS 3' incomplete TSL:5
Ppp6c-203	ENSMUST00000204257.2	1358	<u>81aa</u>	Nonsense mediated decay	757	A0A0N4SVE2	TSL:5
Ppp6c-202	ENSMUST00000143733.1	487	No protein	Processed transcript	(25)	-	TSL:3

The strategy is based on the design of *Ppp6c-201* transcript, the transcription is shown below:



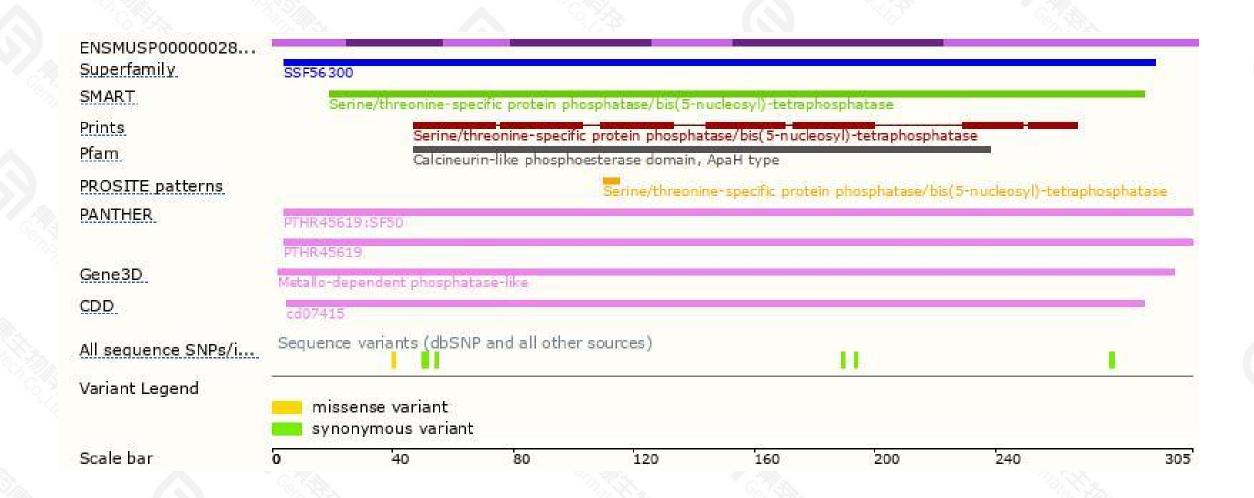
Genomic location distribution





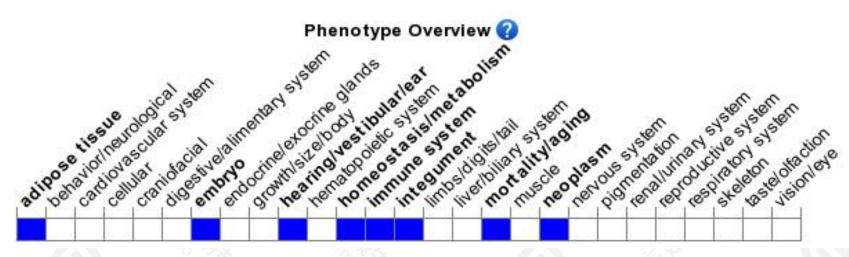
Protein domain





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





