

Prmt6 Cas9-KO Strategy

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Project Overview



Project Name Prmt6

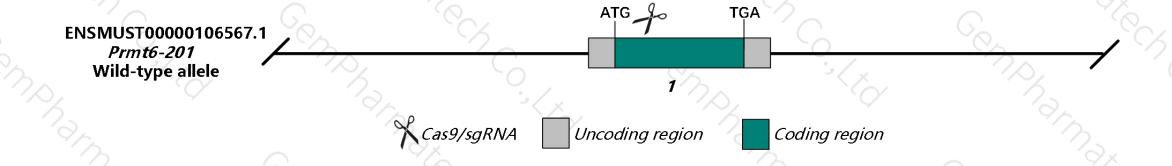
Project type Cas9-KO

Strain background C57BL/6N

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Prmt6* gene. The schematic diagram is as follows:



Technical routes



➤ In this project we use CRISPR/Cas9 technology to modify *Prmt6* 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 *Prmt6* gene is located on the Chr3. 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)



Prmt6 protein arginine N-methyltransferase 6 [Mus musculus (house mouse)]

Gene ID: 99890, updated on 27-Aug-2019





Official Symbol Prmt6 provided by MGI

Official Full Name protein arginine N-methyltransferase 6 provided by MGI

Primary source MGI:MGI:2139971

See related Ensembl:ENSMUSG00000049300

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 Hrmt1l6; AW124876; BB233495

Expression Ubiquitous expression in ovary adult (RPKM 8.9), adrenal adult (RPKM 6.9) and 28 other tissues See more

Orthologs human all

Transcript information (Ensembl)

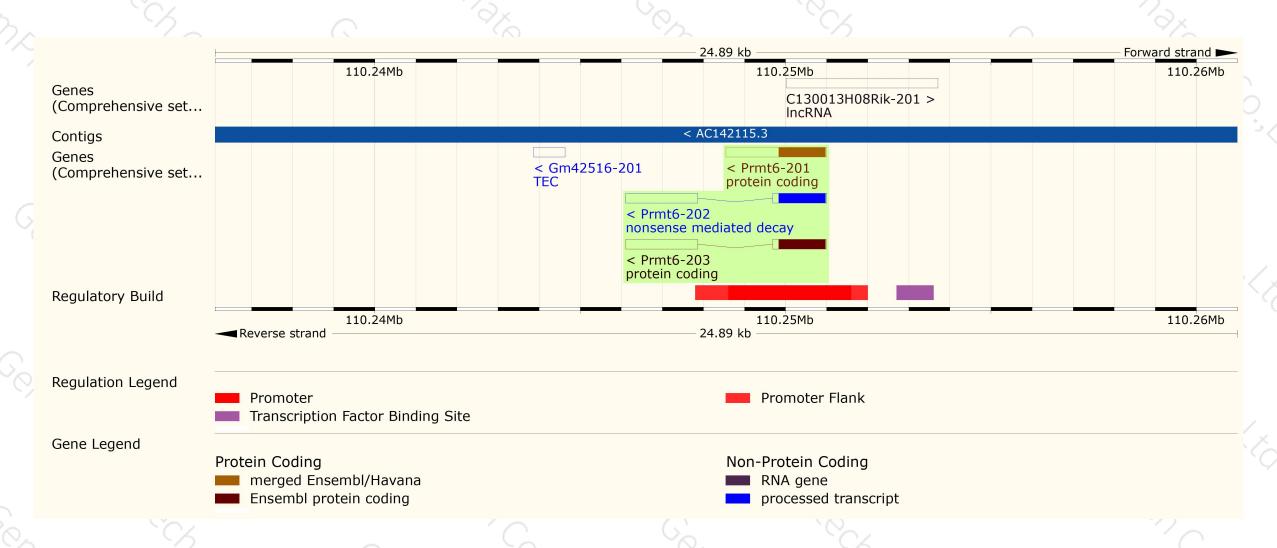


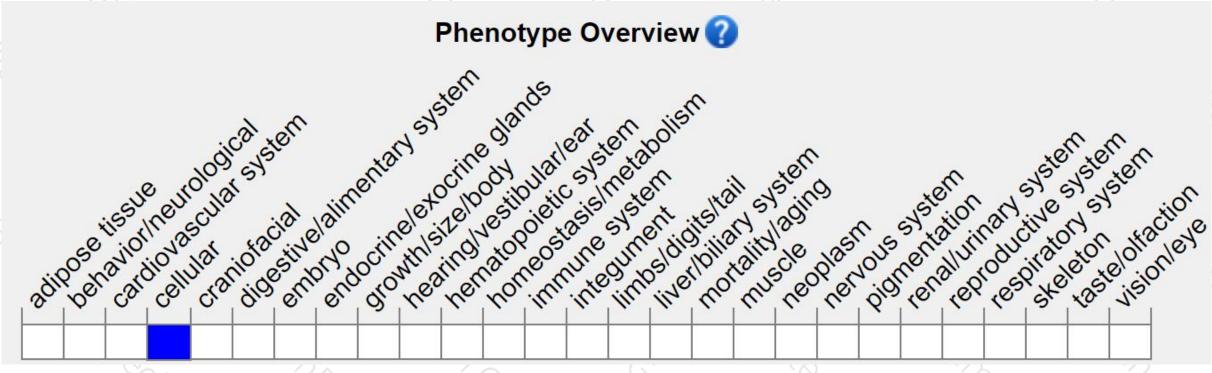
The gene has 3 transcripts, all transcripts are shown below:

Name 🍦	Transcript ID 🛊	bp 🌲	Protein	Translation ID 🔷	Biotype	CCDS	UniProt 🍦	Flags		
Prmt6-203	ENSMUST00000190378.1	3051	<u>378aa</u>	ENSMUSP00000140836.1	Protein coding	CCDS38605 ₽	Q6NZB1₽	TSL:1	GENCODE basic	APPRIS P1
Prmt6-201	ENSMI Stable ID 06567.1	2450	<u>378aa</u>	ENSMUSP00000102177.1	Protein coding	CCDS38605 ₺	Q6NZB1₺	TSL:NA	GENCODE basic	APPRIS P1
Prmt6-202	ENSMUST00000168412.2	3059	<u>378aa</u>	ENSMUSP00000129801.1	Nonsense mediated decay	CCDS38605 ₽	Q6NZB1₽		TSL:1	

Genomic location distribution







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

Mice homozygous for a knock-out allele exhibit decreased mouse embryonic fibroblast proliferation and early cellular replicative senescence.



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

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