

Pde8a Cas9-KO Strategy

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Project Overview



Project Name

Pde8a

Project type

Cas9-KO

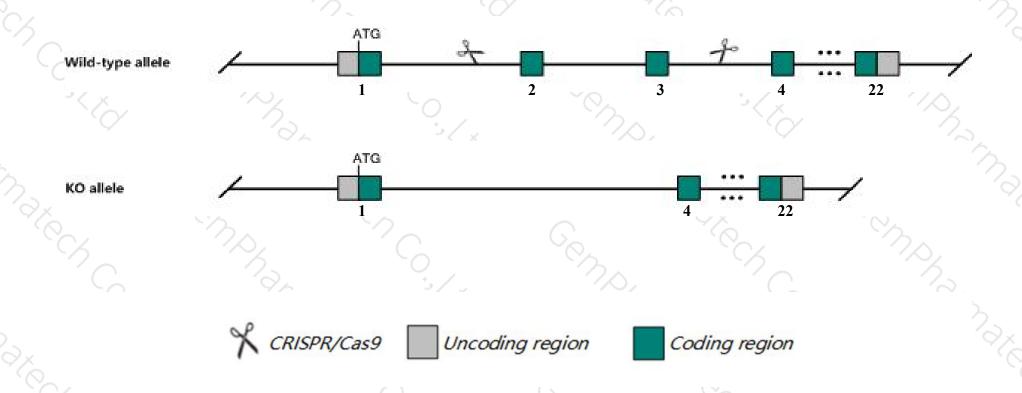
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pde8a* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Pde8a* gene has 3 transcripts. According to the structure of *Pde8a* gene, exon2-exon3 of *Pde8a-201*(ENSMUST00000026672.7) transcript is recommended as the knockout region. The region contains 248bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Pde8a* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ According to the existing MGI data, Targeted disruption of this gene results in a 4-fold increase in basal release of testosterone in isolated Leydig cells as well as a significant increase in the sensitivity to luteinizing hormone, measured as testosterone released into the media.
- > The *Pde8a* gene is located on the Chr7. 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)



Pde8a phosphodiesterase 8A [Mus musculus (house mouse)]

Gene ID: 18584, updated on 31-Jan-2019

Summary

↑ ?

Official Symbol Pde8a provided by MGI

Official Full Name phosphodiesterase 8A provided by MGI

Primary source MGI:MGI:1277116

See related Ensembl:ENSMUSG00000025584

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 Al551852, Pde8

Expression Broad expression in testis adult (RPKM 48.2), kidney adult (RPKM 13.7) and 18 other tissuesSee more

Orthologs human all

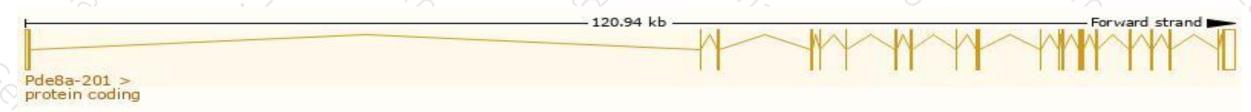
Transcript information (Ensembl)



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

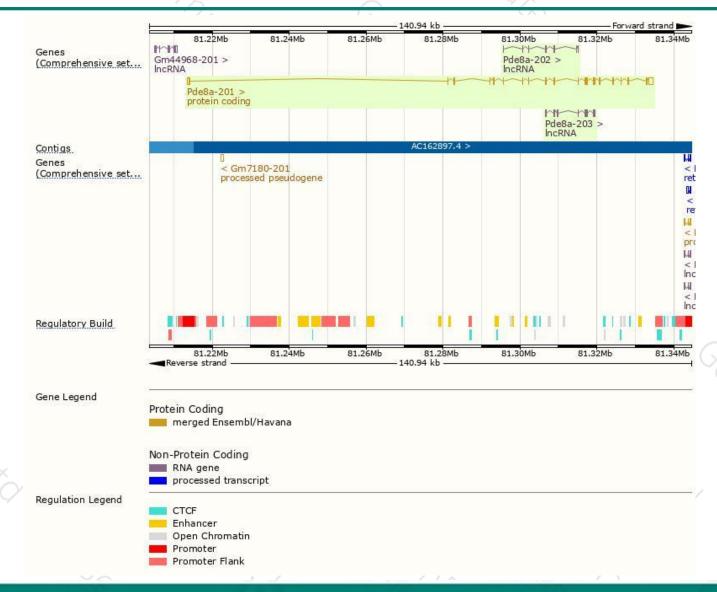
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pde8a-201	ENSMUST00000026672.7	3802	823aa	Protein coding	CCDS40005	<u>088502</u>	TSL:1 GENCODE basic APPRIS P1
Pde8a-203	ENSMUST00000130494.1	861	No protein	IncRNA	-	37.3	TSL:3
Pde8a-202	ENSMUST00000128154.7	583	No protein	IncRNA	¥		TSL:3

The strategy is based on the design of *Pde8a-201* transcript, The transcription is shown below



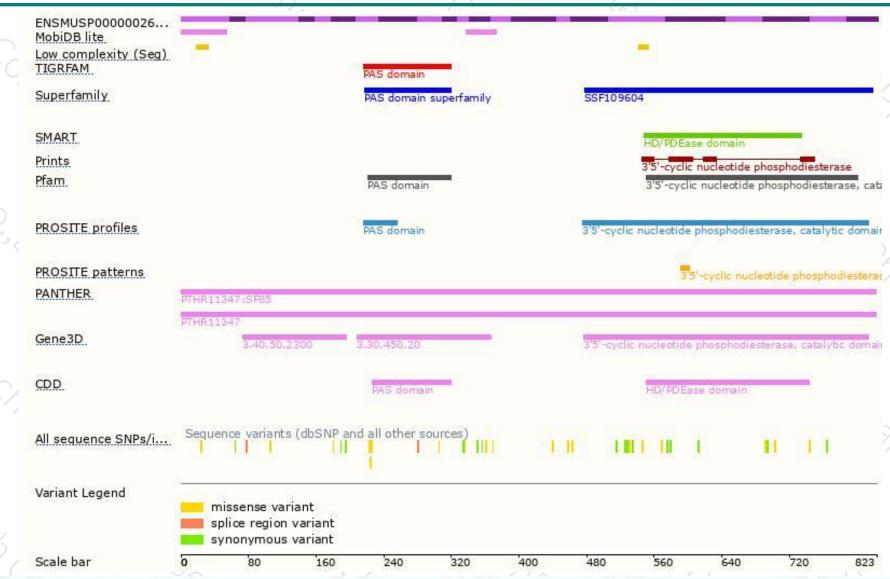
Genomic location distribution





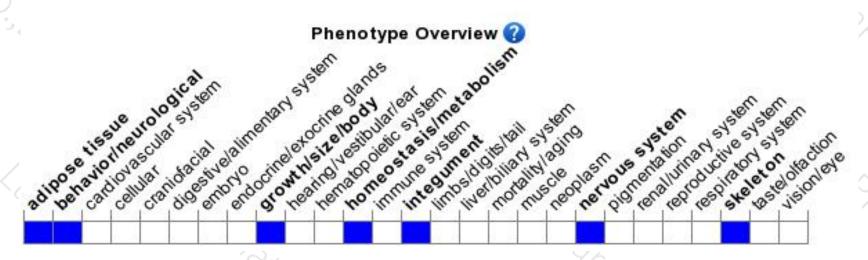
Protein domain





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, Targeted disruption of this gene results in a 4-fold increase in basal release of testosterone in isolated Leydig cells as well as a significant increase in the sensitivity to luteinizing hormone, measured as testosterone released into the media.



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





