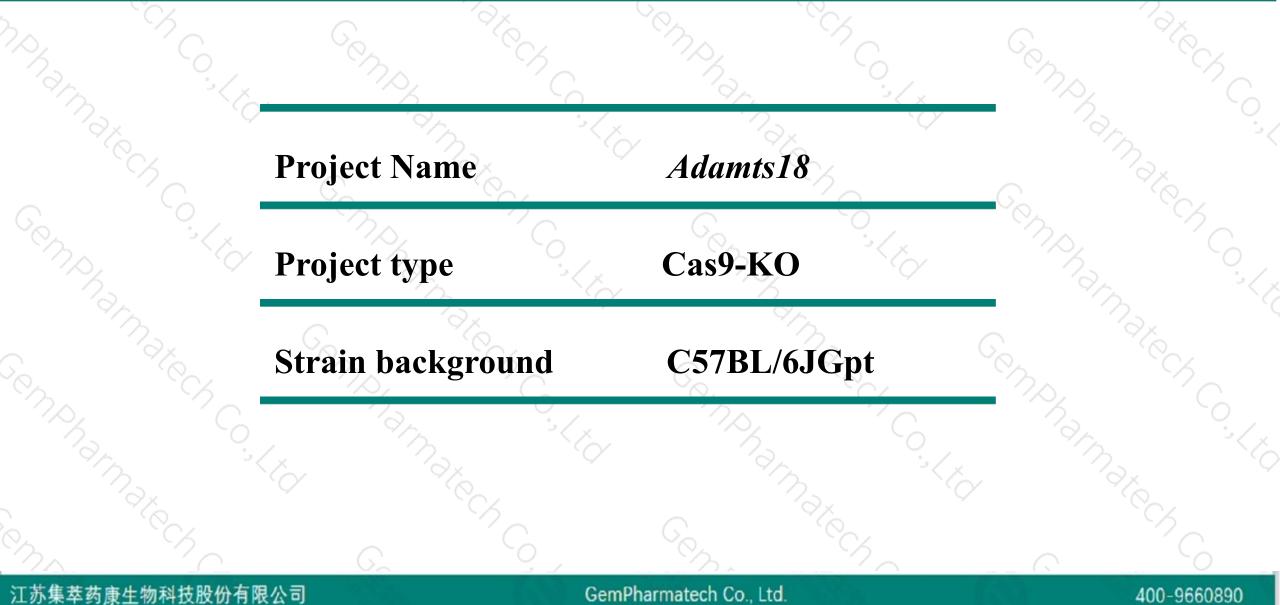


Adamts18 Cas9-KO Strategy

Designer:Xueting Zhang Reviewer:Yanhua Shen Date:2020-02-12

Project Overview

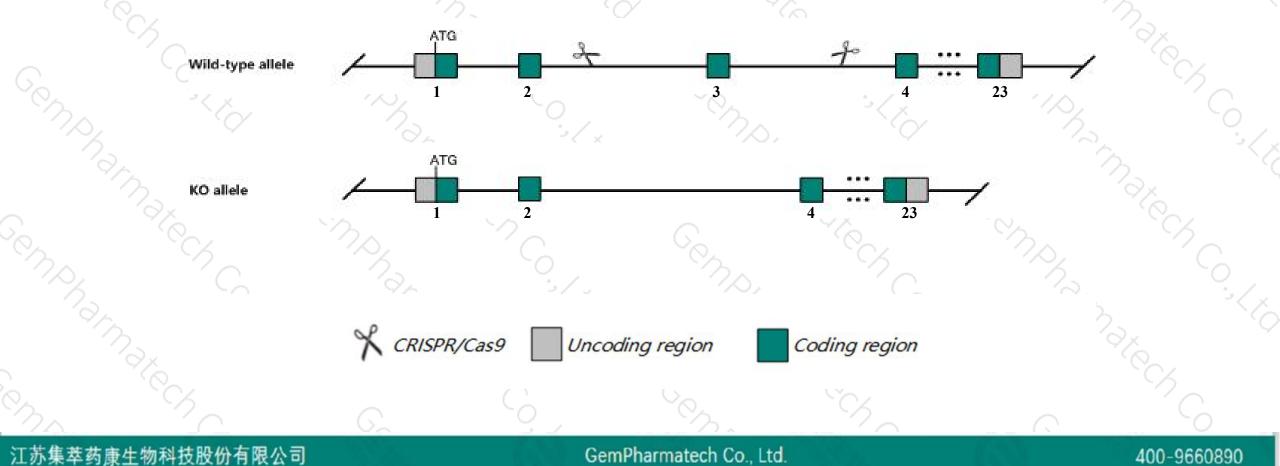




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Adamts18* gene. The schematic diagram is as follows:





- The Adamts18 gene has 7 transcripts. According to the structure of Adamts18 gene, exon3 of Adamts18-201 (ENSMUST00000093113.4) transcript is recommended as the knockout region. The region contains 314bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Adamts18 gene. The brief process is as follows: CRISPR/Cas9 sys

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- According to the existing MGI data, Mice homozygous for a floxed allele exhibit some fertility defects. Mice homozygous for a null allele exhibit growth and eye defects and increased susceptibility to chemically induced tumors.
- > The *Adamts18* gene is located on the Chr8. 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.

Notice

Gene information (NCBI)



Adamts18 a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 18 [*Mus musculus* (house mouse)]

Gene ID: 208936, updated on 24-Dec-2019

Summary

Official Symbol	Adamts18 provided by MGI
Official Full Name	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 18 provided by MGI
Primary source	MGI:MGI:2442600
See related	Ensembl:ENSMUSG00000053399
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	ADAMTS21; 9630038L21; E130314N14Rik
Expression	Biased expression in CNS E14 (RPKM 1.9), CNS E18 (RPKM 1.8) and 12 other tissues See more
Orthologs	human all

Genomic context

2 ?

\$?

Location: 8; 8 E1

Exon count: 24

See Adamts18 in Genome Data Viewer

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	8	NC_000074.6 (113697123113849343, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	8	NC_000074.5 (116222037116372739, complement)

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Transcript information (Ensembl)



The gene has 7 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adamts18-201	ENSMUST0000093113.4	5642	<u>1219aa</u>	Protein coding	CCDS40485	Q4VC17	TSL:1 GENCODE basic APPRIS P1
Adamts18-204	ENSMUST00000212665.1	4591	<u>117aa</u>	Nonsense mediated decay	-8	A0A1D5RLE1	CDS 5' incomplete TSL:1
Adamts18-202	ENSMUST00000212437.1	3307	<u>148aa</u>	Nonsense mediated decay	-	A0A1D5RLA5	CDS 5' incomplete TSL:1
Adamts18-207	ENSMUST00000213078.1	3524	No protein	Retained intron	<u>-</u> 2	12	TSL:1
Adamts18-205	ENSMUST00000213061.1	3397	No protein	Retained intron	5	17	TSL:1
Adamts18-206	ENSMUST00000213076.1	693	No protein	IncRNA	-8	8	TSL:3
Adamts18-203	ENSMUST00000212527.1	637	No protein	IncRNA	2	-	TSL:3

The strategy is based on the design of Adamts18-201 transcript, The transcription is shown below

< Adamts18-201 protein coding

Reverse strand -

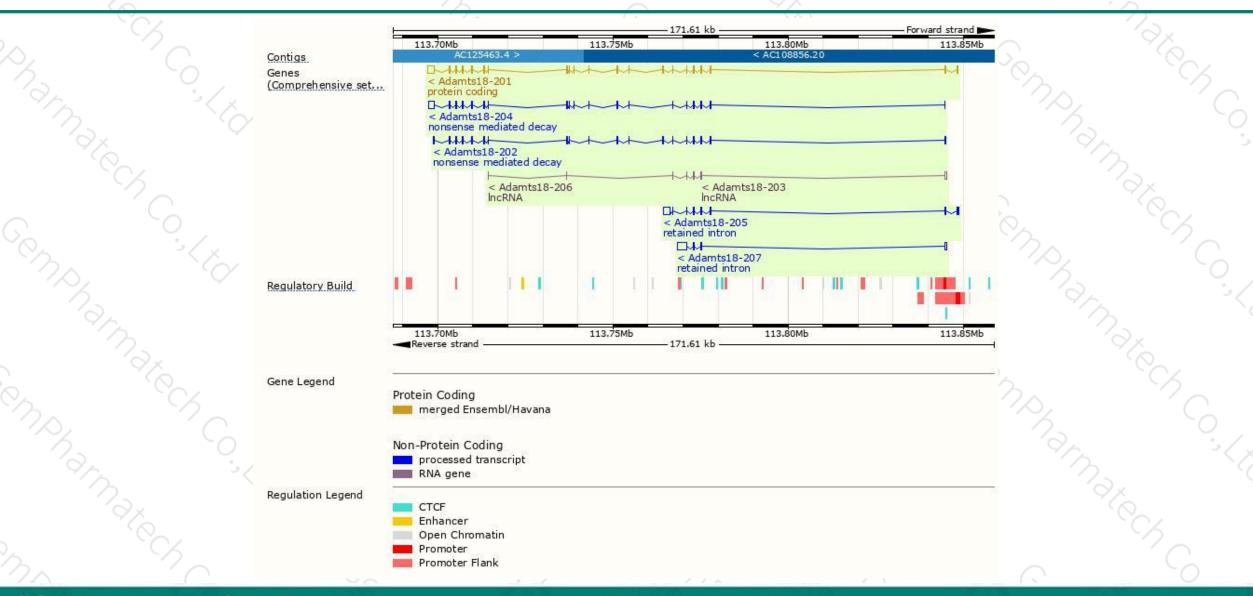
- 151.49 kb -

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Genomic location distribution



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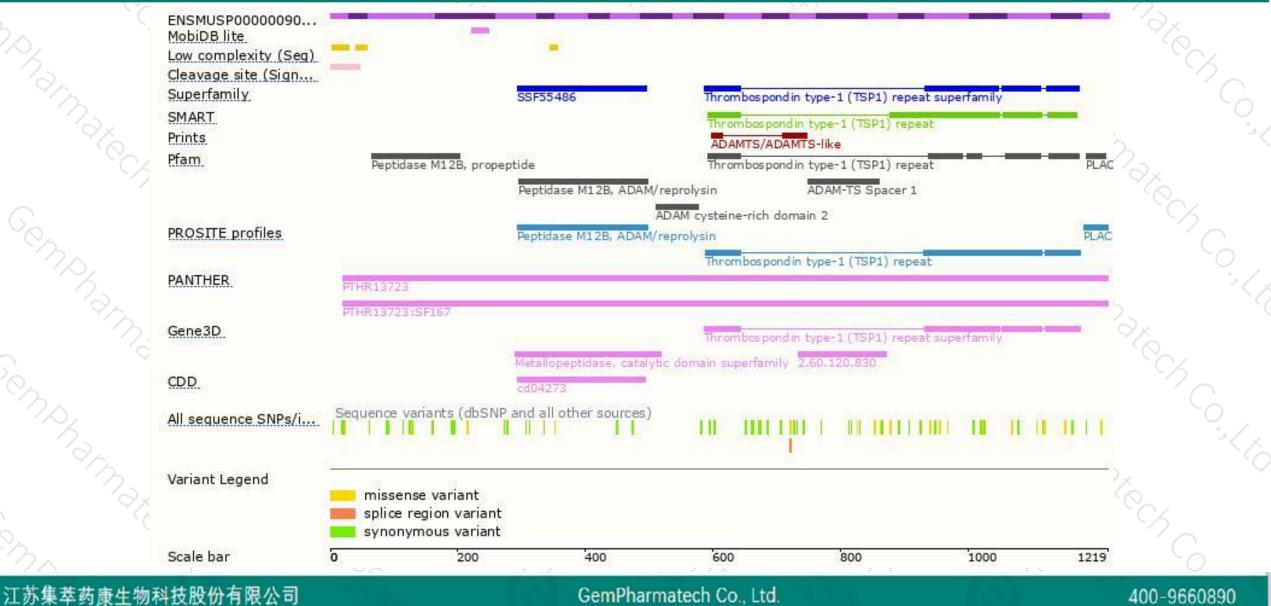
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Protein domain





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, Mice homozygous for a floxed allele exhibit some fertility defects. Mice homozygous for a null allele exhibit growth and eye defects and increased susceptibility to chemically induced tumors.



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



