

Gramd4 Cas9-KO Strategy

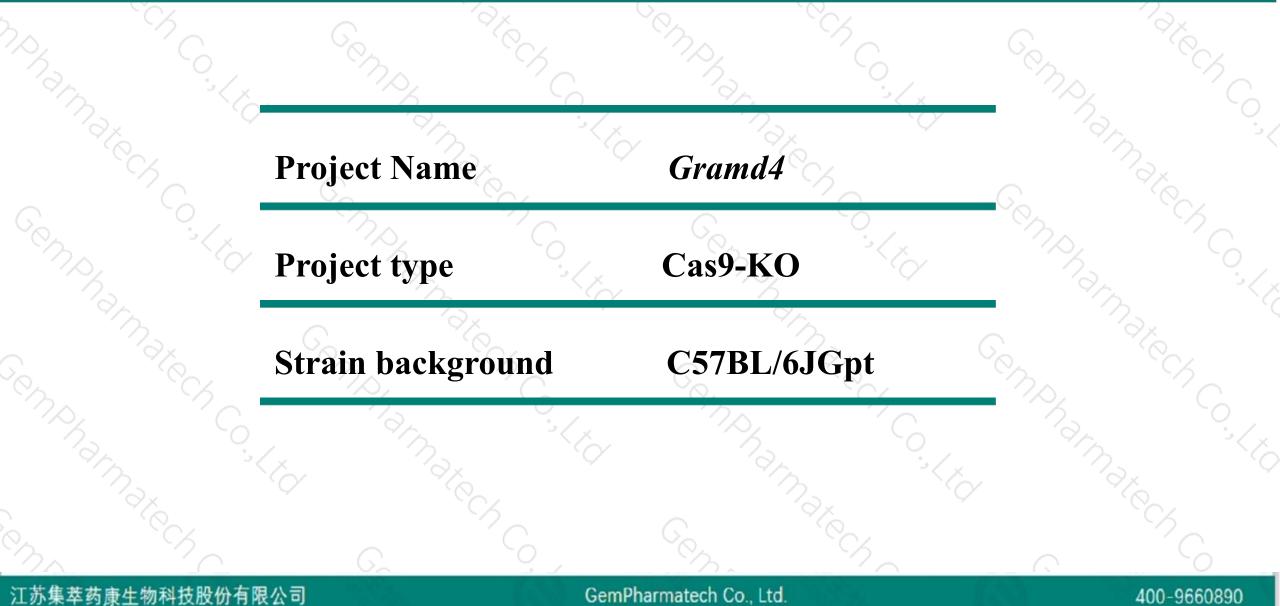
Designer: Xueting Zhang

Reviewer: Daohua Xu

Design Date: 2020-8-11

Project Overview

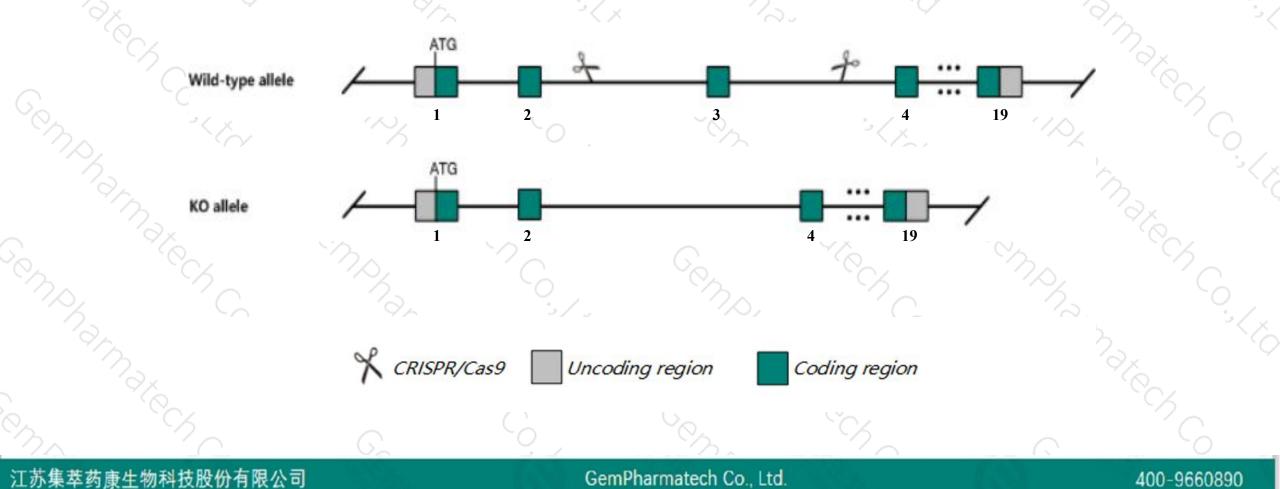




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gramd4* gene. The schematic diagram is as follows:





➤ The Gramd4 gene has 5 transcripts. According to the structure of Gramd4 gene, exon3 of Gramd4-201(ENSMUST00000088931.9) transcript is recommended as the knockout region. The region contains 121bp coding sequence. Knock out the region will result in disruption of protein function.

➤ In this project we use CRISPR/Cas9 technology to modify *Gramd4* 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.

> The *Gramd4* 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.

► Transcript *Gramd4*-205 may not be affected.

> 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)



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Gramd4 GRAM domain containing 4 [Mus musculus (house mouse)]

Gene ID: 223752, updated on 13-Mar-2020

Summary

Official Symbol	Gramd4 provided by MGI
Official Full Name	GRAM domain containing 4 provided by MGI
Primary source	MGI:MGI:2676308
See related	Ensembl:ENSMUSG0000035900
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	9930016013
Expression	Ubiquitous expression in whole brain E14.5 (RPKM 10.5), CNS E18 (RPKM 10.3) and 27 other tissuesSee more
Orthologs	human all

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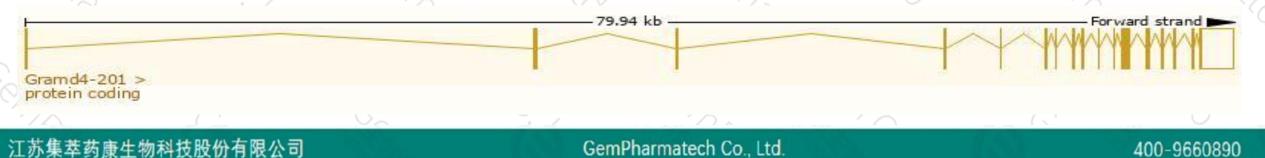
Transcript information (Ensembl)



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

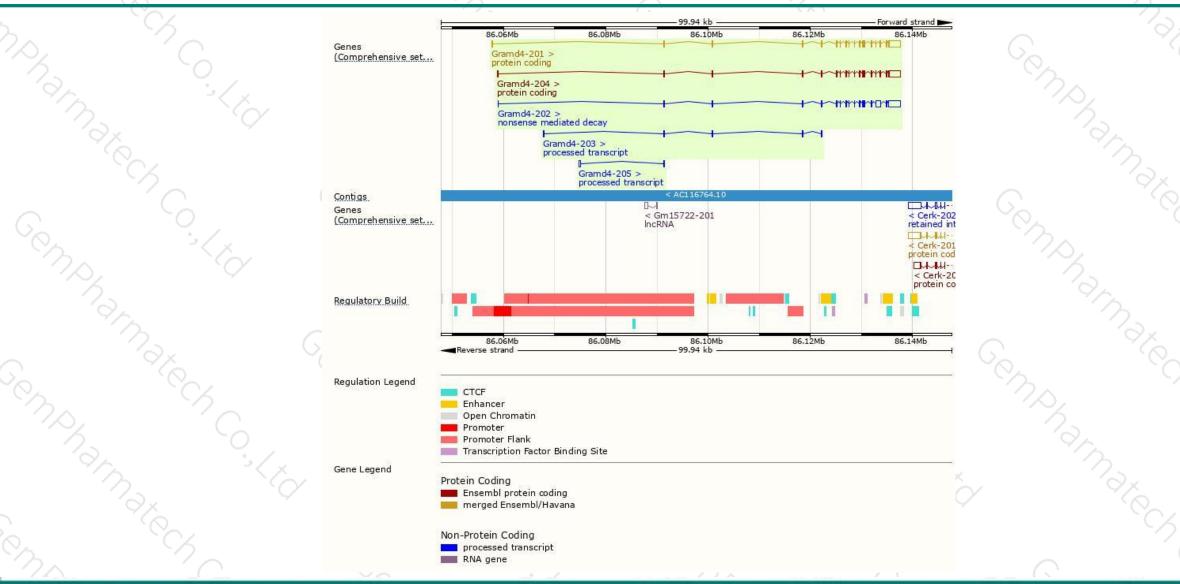
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags		
Gramd4-204	ENSMUST00000138134.7	4292	<u>627aa</u>	Protein coding	CCDS56998	D3YUE7	TSL:1 GENCODE basic		
Gramd4-201	ENSMUST0000088931.9	4115	<u>633aa</u>	Protein coding	CCDS27727	<u>Q8CB44</u>	TSL:1 GENCODE basic APPRIS P1		
Gramd4-202	ENSMUST00000123349.1	4861	<u>323aa</u>	Nonsense mediated decay	120	D6RET7	TSL:2		
Gramd4-203	ENSMUST00000123474.1	597	No protein	Processed transcript	670		TSL:5		
Gramd4-205	ENSMUST00000147286.1	374	No protein	Processed transcript	343	12	TSL:2		

The strategy is based on the design of *Gramd4-201* transcript, the transcription is shown below:



Genomic location distribution





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Protein domain



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



