

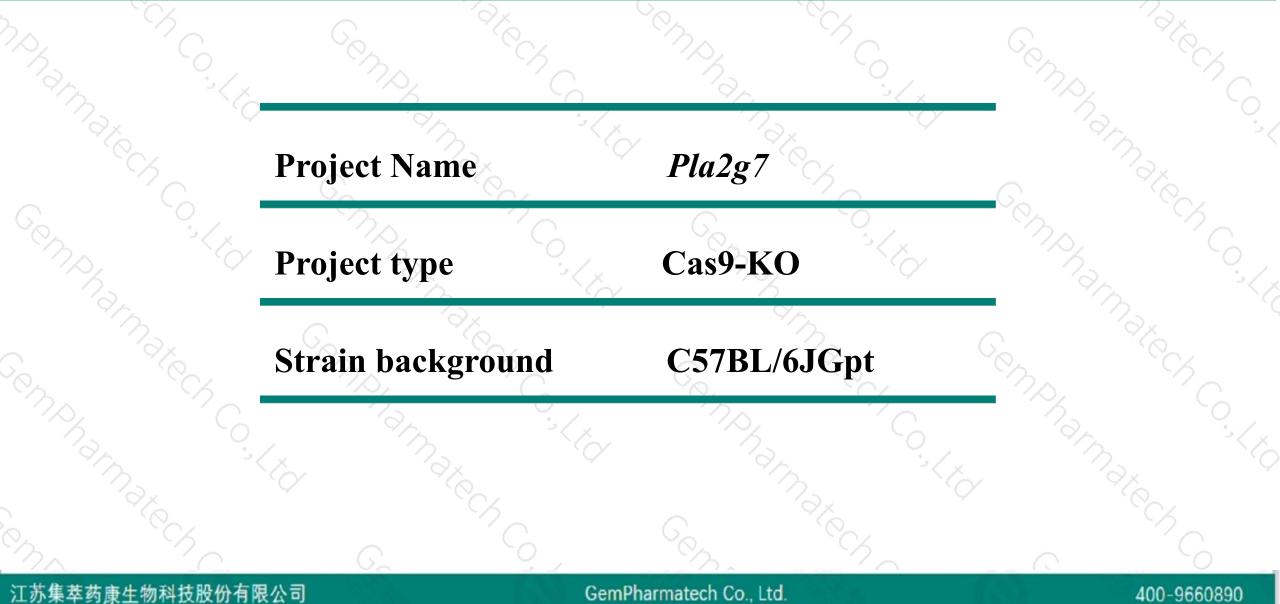
# Pla2g7 Cas9-KO Strategy

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### **Project Overview**

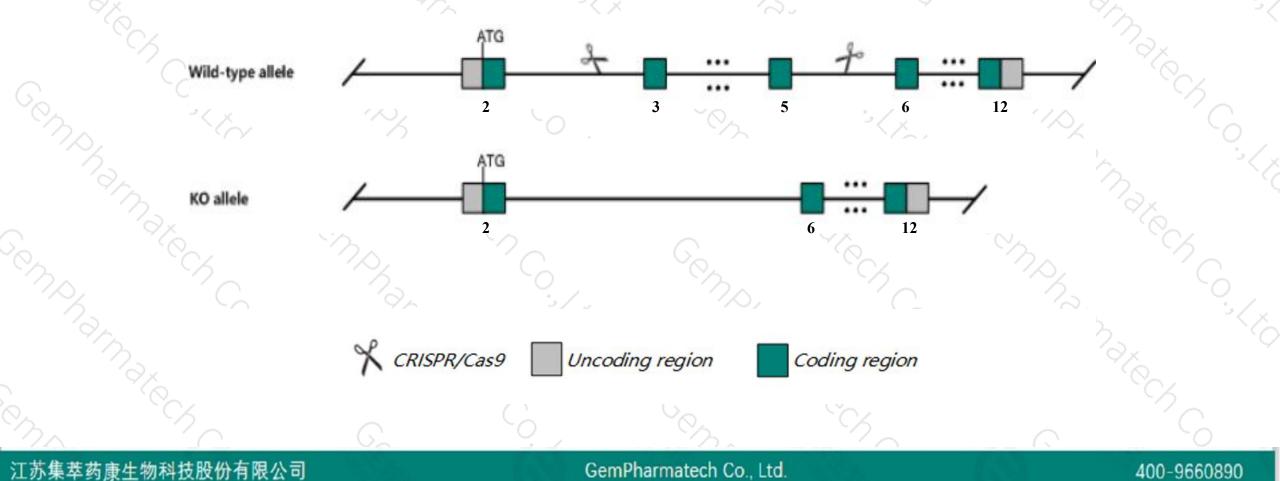




# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Pla2g7* gene. The schematic diagram is as follows:





- The Pla2g7 gene has 6 transcripts. According to the structure of Pla2g7 gene, exon3-exon5 of Pla2g7-201 (ENSMUST0000024706.11) transcript is recommended as the knockout region. The region contains 361bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Pla2g7* gene. The brief process is as follows: CRISPR/Cas9 system



- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit reduced early mortality in response to bacterial exposure, formula feeding and asphyxia, but survivors show a significantly higher incidence of necrotizing enterocolitis relative to wild-type controls.
- ➤ Transcript *Pla2g7*-203 may not be affected.
- The knockout region is near to the N-terminal of 1700071M16Rik gene, this strategy may influence the regulatory function of the N-terminal of 1700071M16Rik gene.
- The Pla2g7 gene is located on the Chr17. 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.

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## Gene information (NCBI)



Pla2g7 phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma) [ Mus musculus (house mouse)] Gene ID: 27226, updated on 12-Aug-2019 Summary \$ ? Official Symbol Pla2g7 provided by MGI Official Full Name phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma) provided by MGI Primary source MGI:MGI:1351327 See related Ensembl:ENSMUSG00000023913 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 PAF-AH; R75400 Expression Broad expression in cerebellum adult (RPKM 43.2), frontal lobe adult (RPKM 30.8) and 16 other tissues See more Orthologs human all Genomic context \$ ? See Pla2g7 in Genome Data Viewer Location: 17; 17 B3 Exon count: 14

Annotation release	Status	Assembly	Chr	Location	2,4
108	current	GRCm38.p6 (GCF_000001635.26)	17	NC_000083.6 (4356799443612202)	1
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	17	NC_000083.5 (4370540043749150)	

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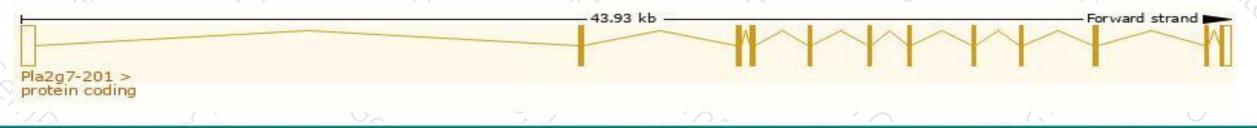
## **Transcript information (Ensembl)**



#### The gene has 6 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pla2g7-201	ENSMUST00000024706.11	2097	<u>440aa</u>	Protein coding	CCDS28796	<u>Q60963</u>	TSL:1 GENCODE basic APPRIS P1
Pla2g7-206	ENSMUST00000169694.1	1112	<u>298aa</u>	Protein coding	-	E9Q330	CDS 3' incomplete TSL:1
Pla2g7-204	ENSMUST00000167214.7	740	<u>211aa</u>	Protein coding	<b>1</b> 0	E9Q6J0	CDS 3' incomplete TSL:5
Pla2g7-205	ENSMUST00000167418.7	551	<u>122aa</u>	Protein coding	20	E9Q4T5	CDS 3' incomplete TSL:3
Pla2g7-202	ENSMUST00000163489.1	1374	No protein	Retained intron	5	-	TSL:1
Pla2g7-203	ENSMUST00000165706.7	727	No protein	IncRNA	-13		TSL:2

The strategy is based on the design of *Pla2g7-201* transcript, The transcription is shown below

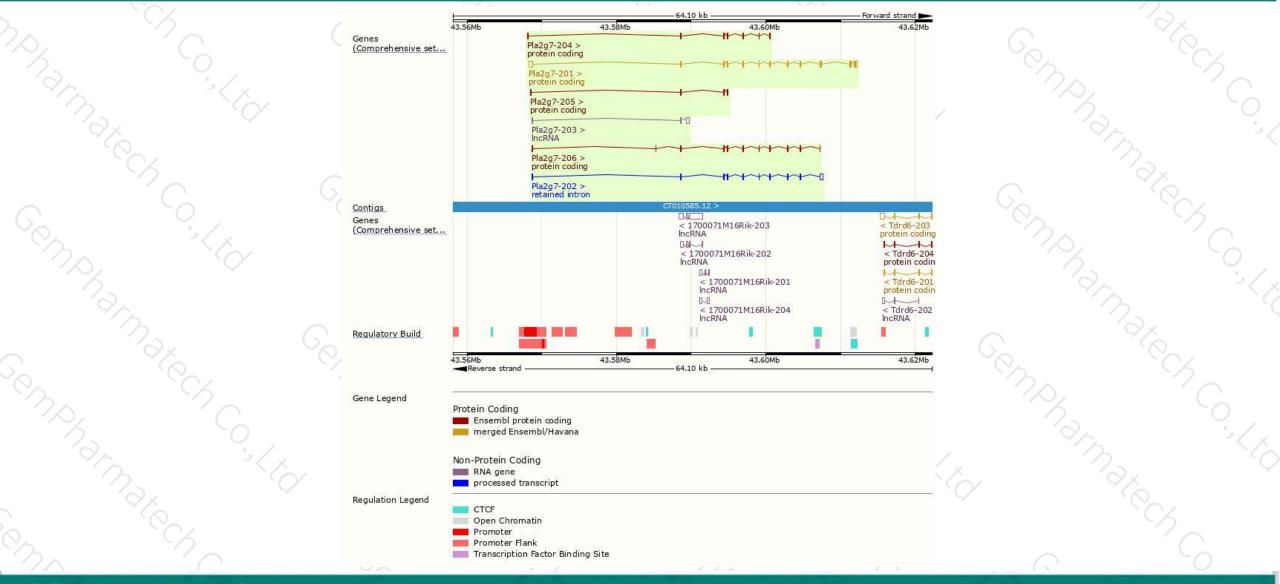


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### **Genomic location distribution**





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### **Protein domain**



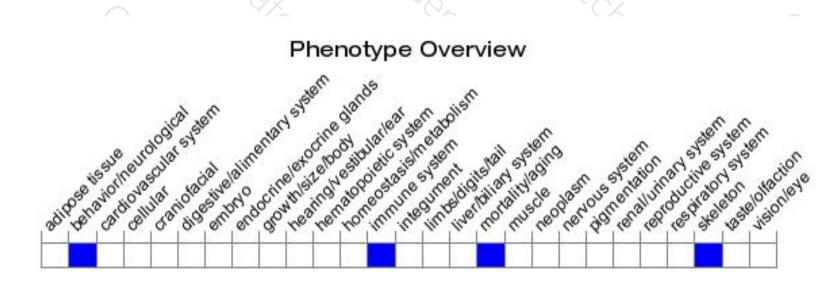
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Yam.	And in case of the second s	elet-activating facto		:-like				
<u>IRSF</u>	Platelet-activating	factor acetylhydrol	ise, eucaryote					
ANTHER	PTHR10272:SF12							1.7
	Platelet-activating	factor acetylhydrol	ise-like					
Sene3D	Alph	a/Beta hydrolase fo	Id					
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### 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 knock-out allele exhibit reduced early mortality in response to bacterial exposure, formula feeding and asphyxia, but survivors show a significantly higher incidence of necrotizing enterocolitis relative to wild-type controls.

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



