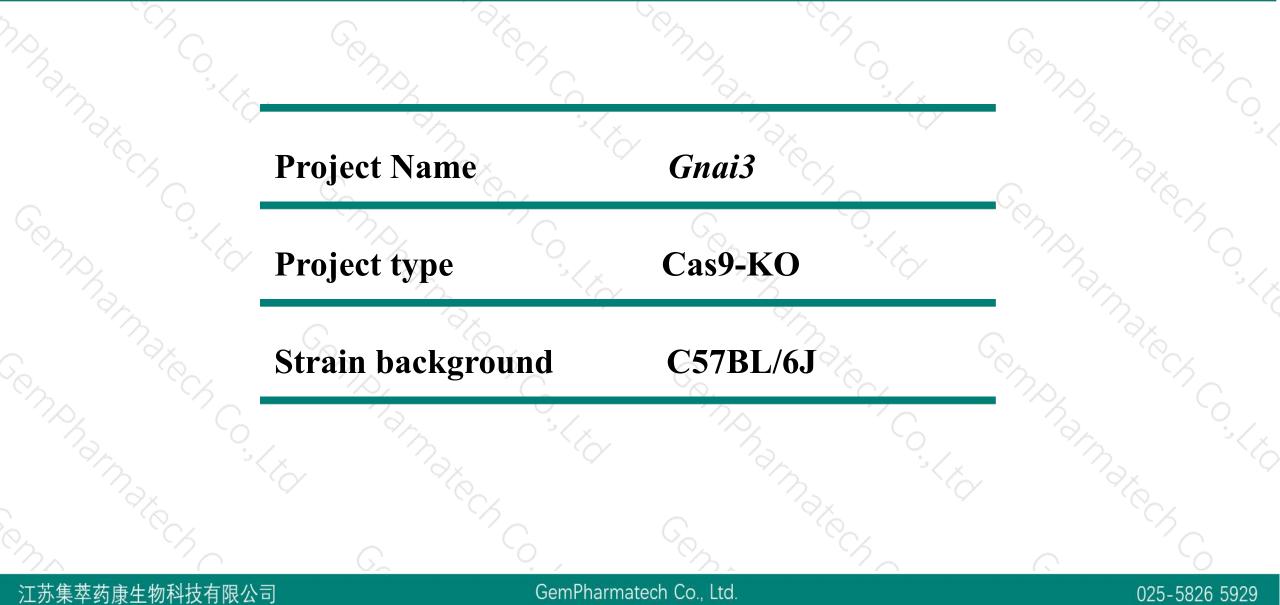


# **Gnai3** Cas9-KO Strategy

Designer: Reviewer: Design Date: JiaYu Xiaojing Li 2019-11-5

### **Project Overview**

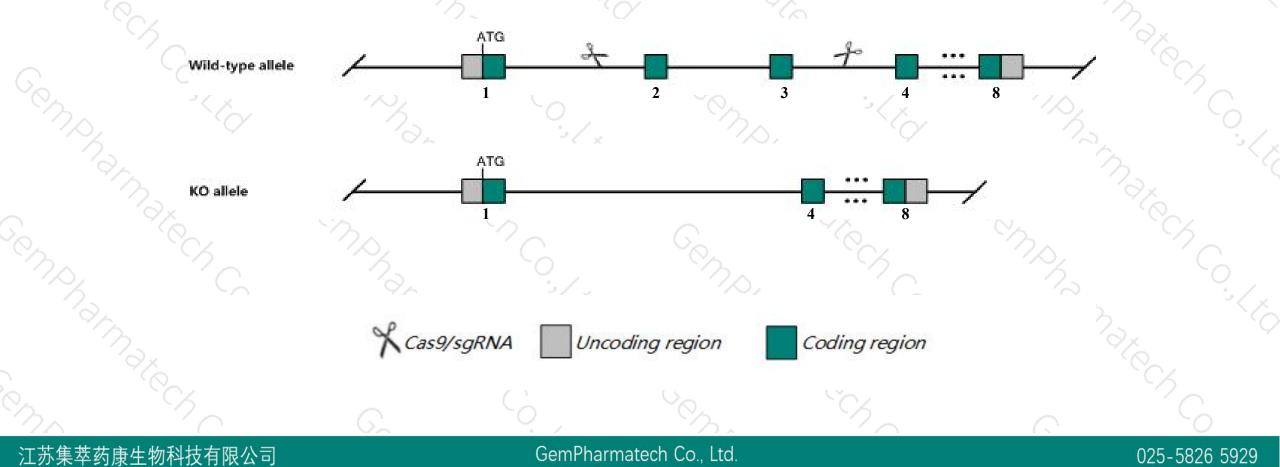




# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Gnai3 gene. The schematic diagram is as follows:





- The Gnai3 gene has 1 transcript. According to the structure of Gnai3 gene, exon2-exon3 of Gnai3-201 (ENSMUST0000000001.4) transcript is recommended as the knockout region. The region contains 185bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gnai3* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

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- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit normal basal cardiac function and beta-adrenergic sensitivity. Mice homozygous for a different knock-out allele exhibit enhanced T cell migration toward CXCR3 agonists.
- The Gnai3 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.

Notice

# **Gene information** (NCBI)



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### Gnai3 guanine nucleotide binding protein (G protein), alpha inhibiting 3 [Mus musculus (house mouse)]

Gene ID: 14679, updated on 7-Apr-2019

#### Summary

Gnai3 provided by MGI
guanine nucleotide binding protein (G protein), alpha inhibiting 3 provided by MGI
MGI:MGI:95773
Ensembl:ENSMUSG000000001
protein coding
VALIDATED
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
Muroidea; Muridae; Murinae; Mus; Mus
AI158965, AW537698, Galphai3, Gnai-3
Ubiquitous expression in placenta adult (RPKM 67.1), CNS E11.5 (RPKM 42.8) and 26 other tissues See more
human all

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### **Transcript information (Ensembl)**



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The gene has 1 transcript, and the transcript is shown below:

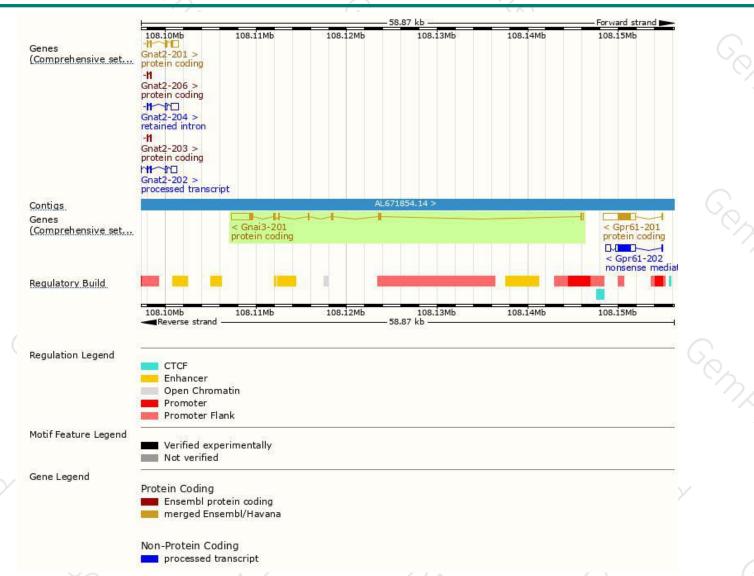
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Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gnai3-201	ENSMUST0000000001.4	3262	<u>354aa</u>	Protein coding	CCDS17751	Q9DC51	TSL:1 GENCODE basic APPRIS P1
23. Maria	K Cont		are ch		Const.	X Co. K.K.	Cemptander (
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Snai3-201 Itein coding	M		N				

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### **Genomic location distribution**





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### **Protein domain**

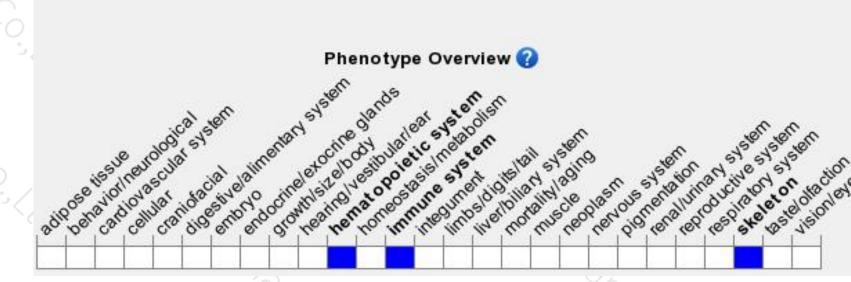


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			G prot	ein alpha s	ubunit, helici	al insertion					
	SMART domains	Guanine n	ucleotide bindi	ing protein	(G-protein),	alpha subunit	- 20 20 20 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10 1	10.2 10.		C	
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	Pfam domain	Guanine r	nucleotide binc	ling protein	(G-protein)	, alpha subunit				10	
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			G prot	ein alpha s	ubunit, helic	al insertion				<u></u>	
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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 normal basal cardiac function and beta-adrenergic sensitivity. Mice homozygous for a different knock-out allele exhibit enhanced T cell migration toward CXCR3 agonists.

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If you have any questions, you are welcome to inquire. Tel: 025-5864 1534



