

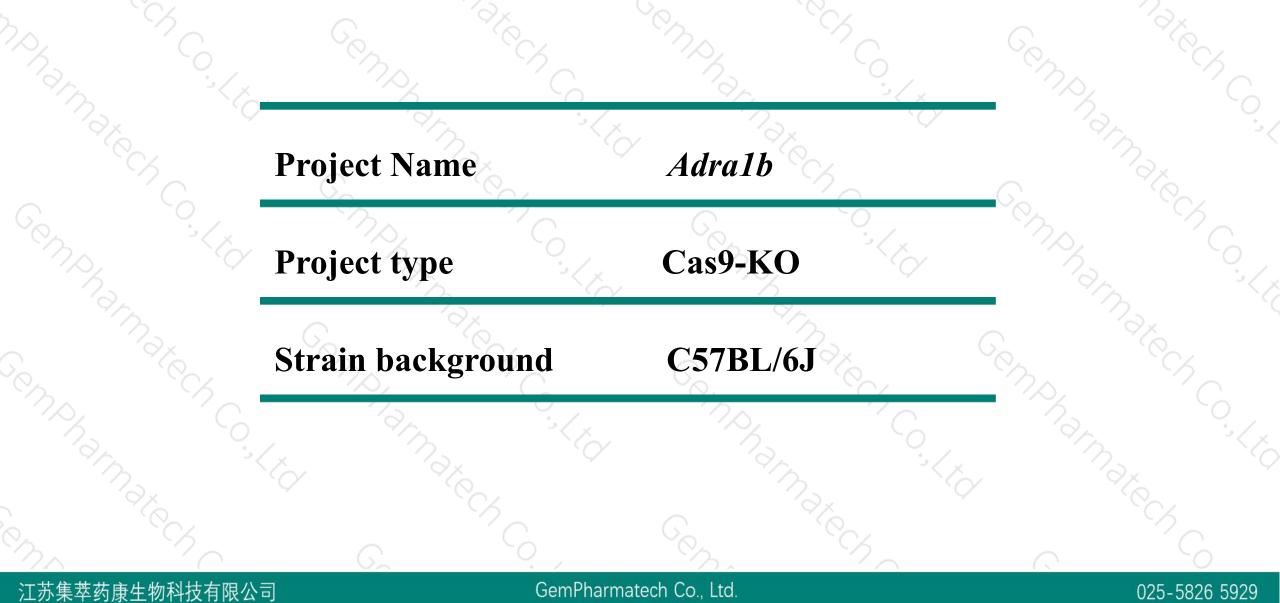
Adra1b Cas9-KO Strategy

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Project Overview





Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Adra1b gene. The schematic diagram is as follows:



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- The Adra1b gene has 4 transcripts. According to the structure of Adra1b gene, exon4 of Adra1b-201 (ENSMUST00000067258.8) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Adra1b* 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, Targeted mutations that inactivate the gene affect atrial contractility and left ventricle function, suggesting their use in modeling chronic heart failure in humans.
- The Adra1b gene is located on the Chr11. 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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Adra1b adrenergic receptor, alpha 1b [Mus musculus (house mouse)]

Gene ID: 11548, updated on 31-Jan-2019

Summary

Official Symbol	Adra1b provided by MGI					
•	adrenergic receptor, alpha 1b provided by MGI					
Primary source	MGI:MGI:104774					
See related	Ensembl:ENSMUSG0000050541					
Gene type	protein coding					
RefSeq status	VALIDATED					
Organism	Mus musculus					
Lineage	Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomo					
	Muroidea; Muridae; Murinae; Mus; Mus					
Also known as	[a]1b					
Expression	Biased expression in liver adult (RPKM 44.6), heart adult (RPKM 12.4) and 6 other tissuesSee more					
Orthologs	human all					



Transcript information (Ensembl)



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The gene has 4 transcripts, all transcripts are shown below:

		T	(C) (1)			T	a kan ta kan
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adra1b-201	ENSMUST0000067258.8	3331	<u>515aa</u>	Protein coding	CCDS24562	Q9DBL0	TSL:1 GENCODE basic APPRIS P1
Adra1b-204	ENSMUST00000167574.1	3047	<u>515aa</u>	Protein coding	CCDS24562	Q9DBL0	TSL:1 GENCODE basic APPRIS P1
Adra1b-203	ENSMUST00000139906.1	1764	<u>462aa</u>	Protein coding	8 <u>4</u>	<u>B1AU41</u>	CDS 3' incomplete TSL:1
Adra1b-202	ENSMUST00000124306.1	3047	No protein	Processed transcript	14	100	TSL:1

The strategy is based on the design of Adra1b-201 transcript, The transcription is shown below

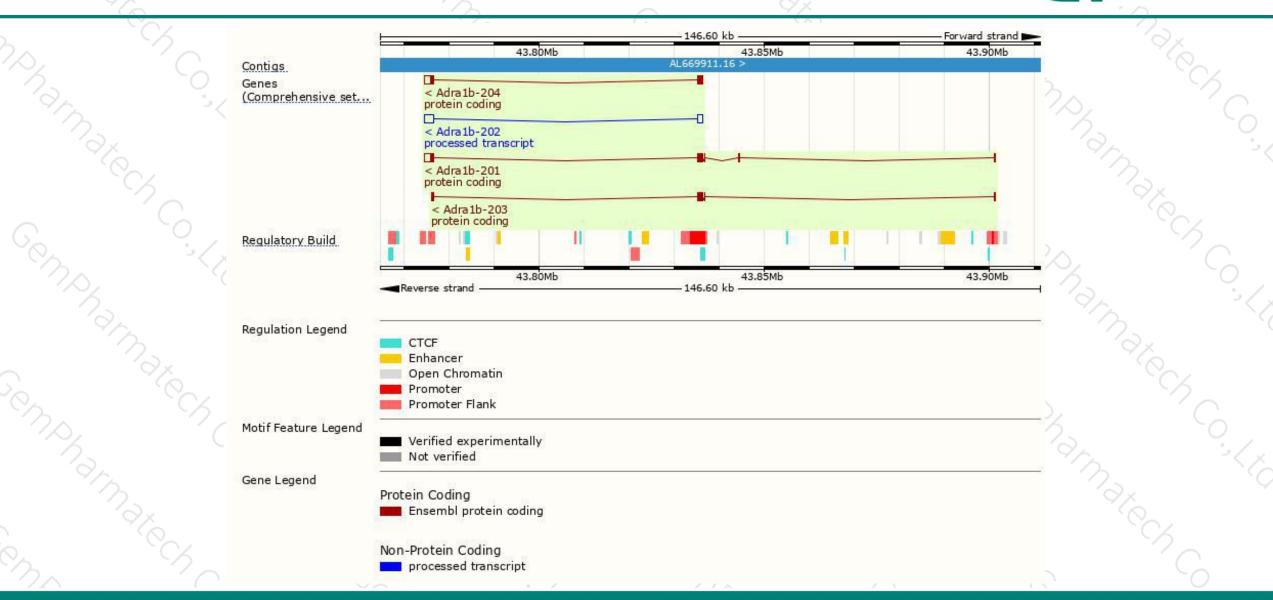
< Adra1b-2	
protein cod	ing

Reverse strand

- 126.61 kb -

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Genomic location distribution



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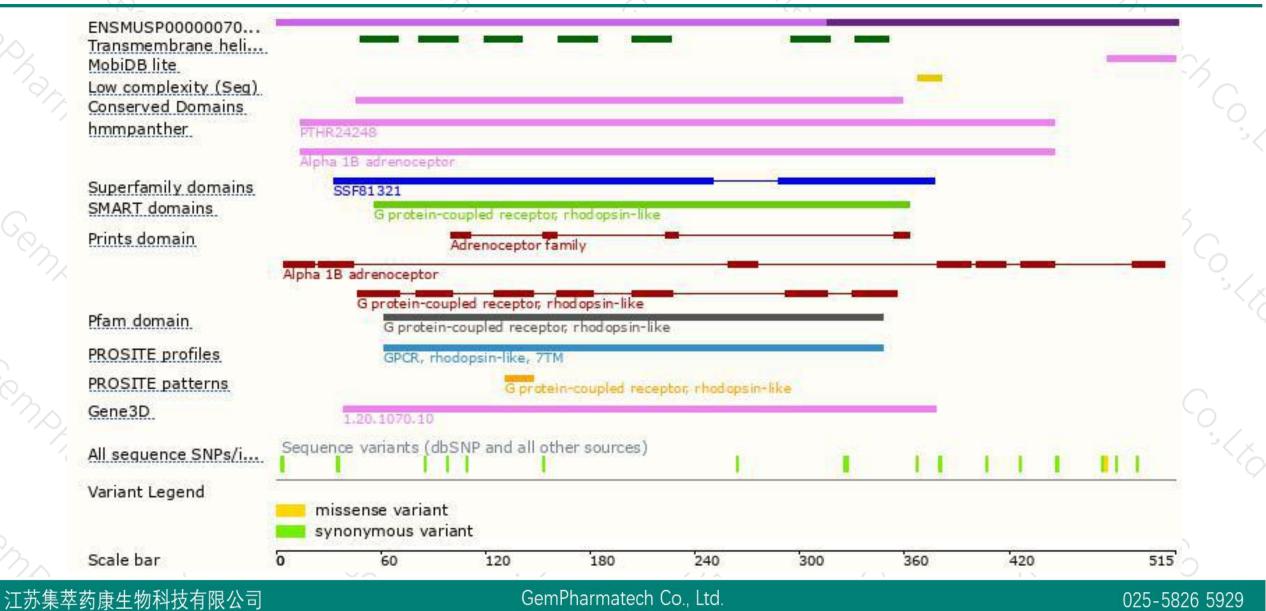
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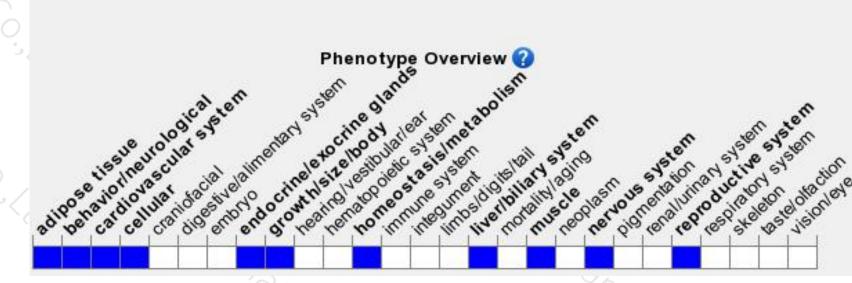
Protein domain





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, Targeted mutations that inactivate the gene affect atrial contractility and left ventricle function, suggesting their use in modeling chronic heart failure in humans.



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



