

# Adra1b Cas9-CKO Strategy

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Reviewer:Fengjuan Wang

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# **Project Overview**



**Project Name** 

Adra1b

**Project type** 

Cas9-CKO

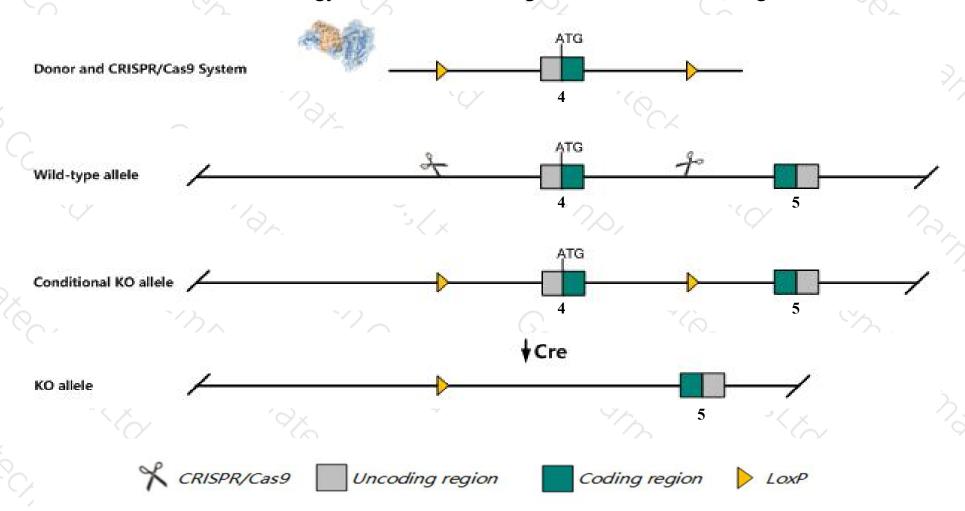
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



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



### Technical routes



- 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:CRISPR/Cas9 system and Donor 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 flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

### **Notice**



- ➤ 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.
- > Since intron 3-4 is only 511bp, the insertion of 5 'loxP may affect the normal splicing of exons.
- The insertion of 5 'loxP into introns upstream of the initial codon may affect potential cis-acting elements and transcription factor binding sites. This strategy removes the original ATG of the gene and may lead to the formation of new unknown proteins
- 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

### Gene information (NCBI)



#### Adra1b adrenergic receptor, alpha 1b [Mus musculus (house mouse)]

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

#### Summary

☆ ?

Official Symbol Adra1b provided by MGI

Official Full Name adrenergic receptor, alpha 1b provided by MGI

Primary source MGI:MGI:104774

See related Ensembl: ENSMUSG00000050541

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 [a]1b

Expression Biased expression in liver adult (RPKM 44.6), heart adult (RPKM 12.4) and 6 other tissuesSee more

Orthologs human all

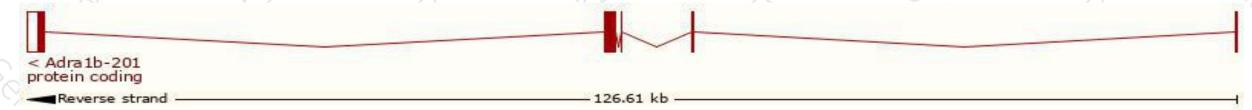
# Transcript information (Ensembl)



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

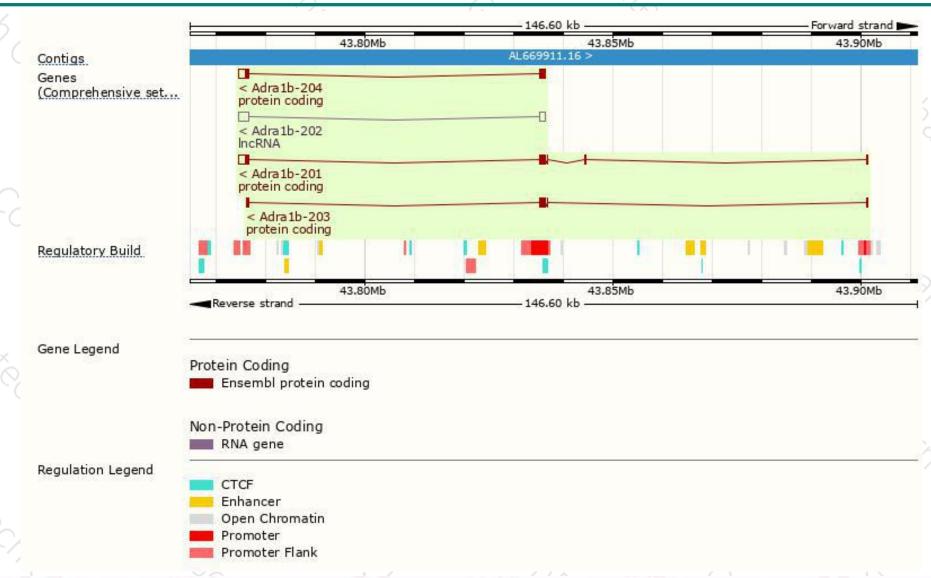
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adra1b-201	ENSMUST00000067258.8	3331	<u>515aa</u>	Protein coding	CCDS24562	Q9DBL0	TSL:1 GENCODE basic APPRIS P1
Adra1b-204	ENSMUST00000167574.1	3047	<u>515aa</u>	Protein coding	CCDS24562	Q9DBL0	TSL:1 GENCODE basic APPRIS P1
Adra1b-203	ENSMUST00000139906.1	1764	<u>462aa</u>	Protein coding	84	B1AU41	CDS 3' incomplete TSL:1
Adra1b-202	ENSMUST00000124306.1	3047	No protein	Processed transcript	(2	323	TSL:1

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



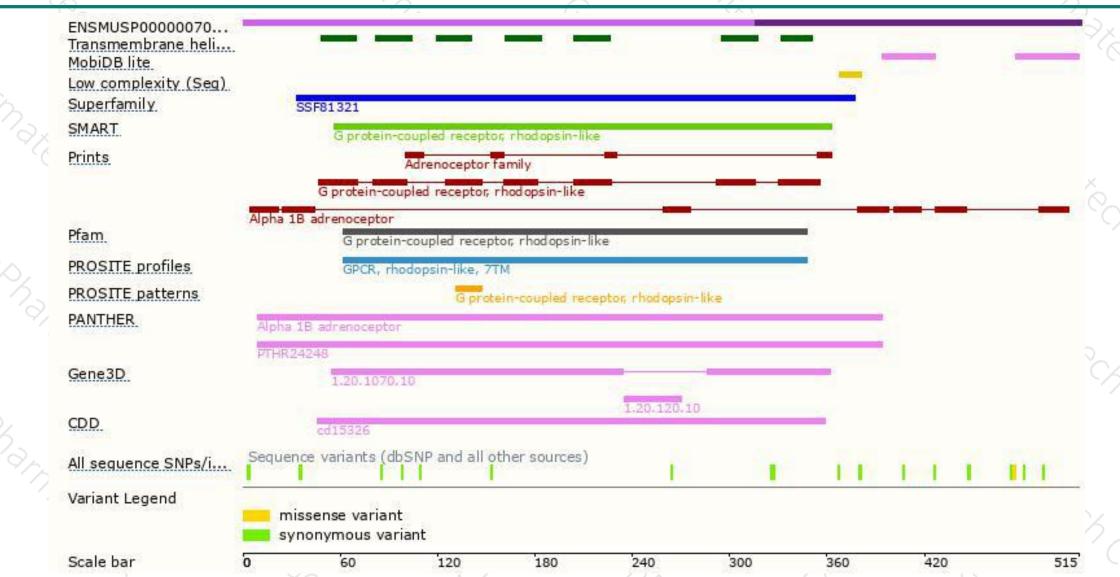
### Genomic location distribution





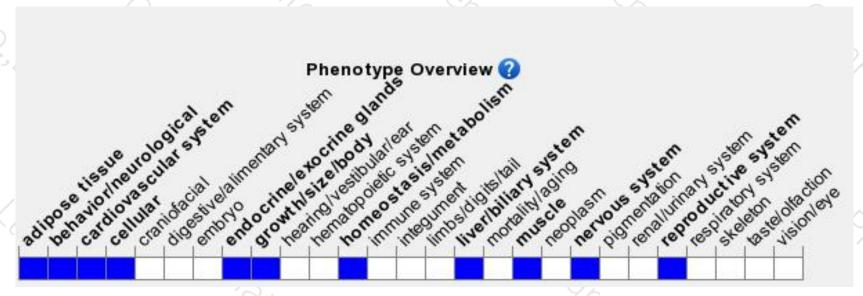
### Protein domain





# 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: 400-9660890





