# *Jph2* Cas9-CKO Strategy

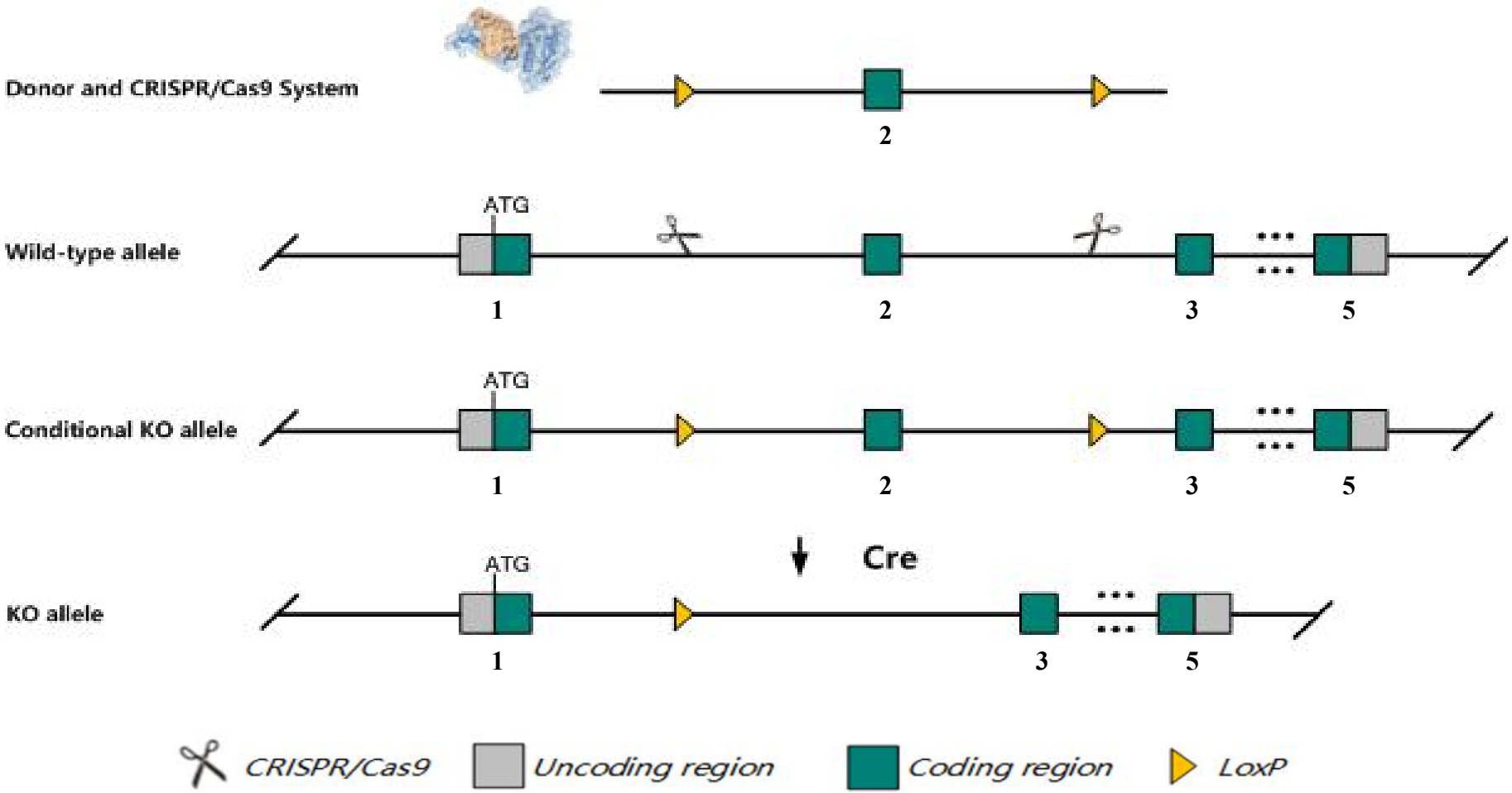
| | |
|---|---|
| **Designer:** | **Huan Fan** |
| **Reviewer** | **Huan Wang** |
| **Design Date:** | **2020-4-16** |

# Project Overview

| | |
|---|---|
| **Project Name** | ***Jph2*** |
| **Project type** | **Cas9-CKO** |
| **Strain background** | **C57BL/6JGpt** |

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Jph2* gene. The schematic diagram is as follows:



# Technical routes

The *Jph2* gene has 2 transcripts. According to the structure of *Jph2* gene, exon2 of *Jph2-201* (ENSMUST00000017961.10) transcript is recommended as the knockout region.The region contains 772bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Jph2* 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,homozygotes for a targeted null mutation exhibit a deficiency of junctional membrane complexes in cardiac myocytes and die by embryonic day 10.5.

The *Jph2* gene is located on the Chr2. 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

集萃药康 GemPharmatech

## Jph2 junctophilin 2 [Mus musculus (house mouse)]

Gene ID: 59091, updated on 13-Mar-2020

### Summary

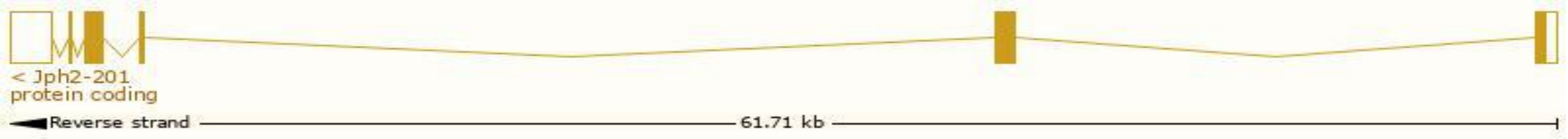
| | |
|---|---|
| **Official Symbol** | Jph2 provided by MGI |
| **Official Full Name** | junctophilin 2 provided by MGI |
| **Primary source** | MGI:MGI:1891496 |
| **See related** | Ensembl:ENSMUSG00000017817 |
| **Gene type** | protein coding |
| **RefSeq status** | VALIDATED |
| **Organism** | Mus musculus |
| **Lineage** | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| **Also known as** | 1110002E14Rik, JP-2, Jp2 |
| **Expression** | Biased expression in heart adult (RPKM 70.1), bladder adult (RPKM 48.2) and 10 other tissues See more |
| **Orthologs** | human all |

# Transcript information Ensembl

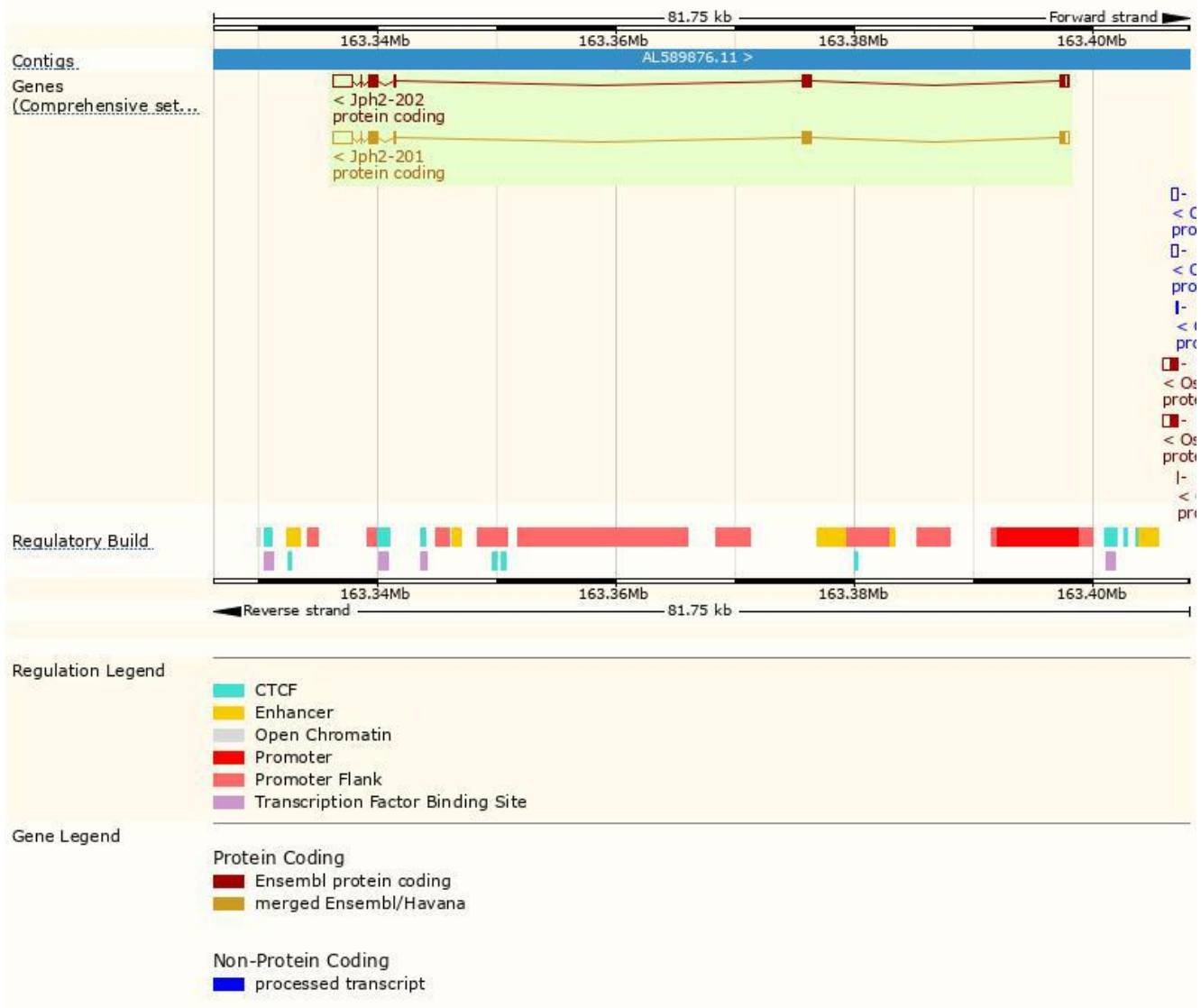
The gene has 2 transcripts,all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|---|---|---|---|---|---|---|---|
| Jph2-201 | ENSMUST00000017961.10 | 4211 | 696aa | Protein coding | CCDS17008 | Q9ET78 | TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1 |
| Jph2-202 | ENSMUST00000109425.2 | 4155 | 696aa | Protein coding | CCDS17008 | Q9ET78 | TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1 |

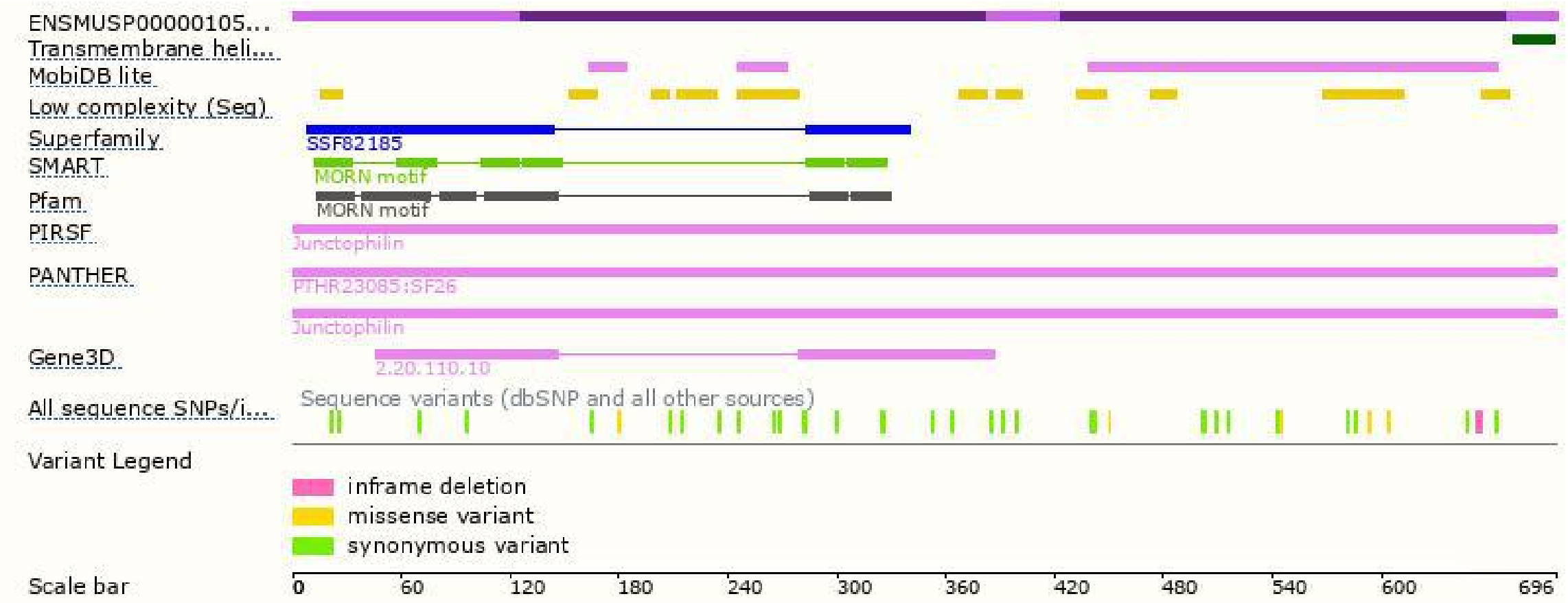
The strategy is based on the design of *Jph2-201* transcript,the transcription is shown below



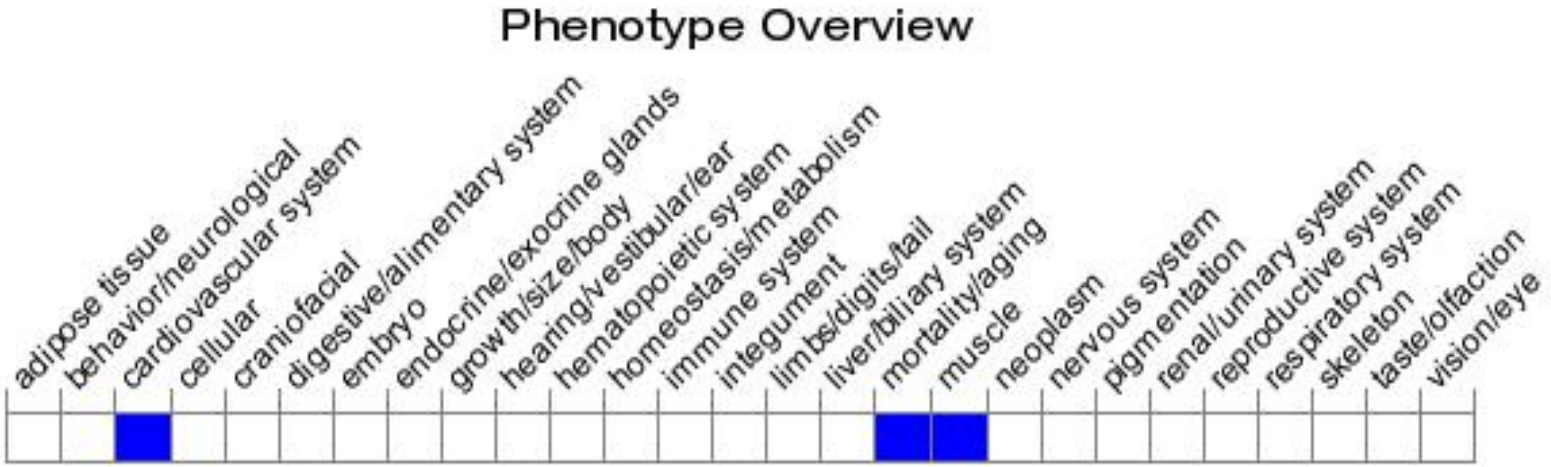
# Genomic location distribution



# 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,homozygotes for a targeted null mutation exhibit a deficiency of junctional membrane complexes in cardiac myocytes and die by embryonic day 10.5.

集萃药康
GemPharmatech

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

集萃药康生物科技
GemPharmatech Co.,Ltd