

# Shc1 Cas9-KO Strategy

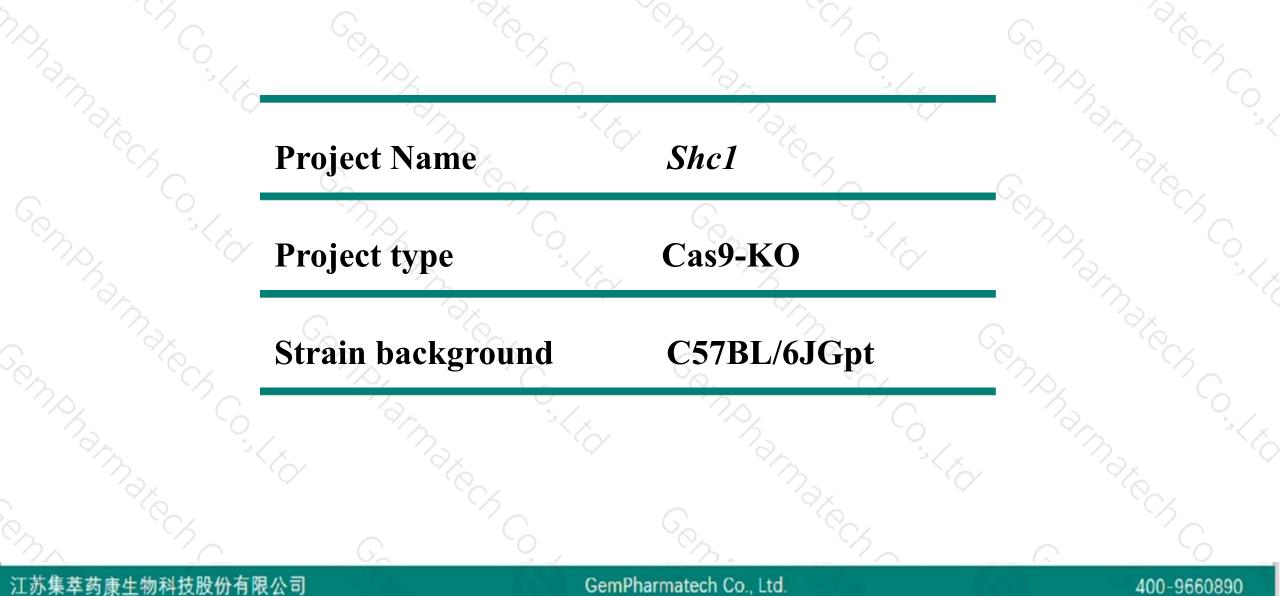
**Designer: Huimin Su** 

**Reviewer: Ruirui Zhang** 

**Design Date: 2020-9-23** 

## **Project Overview**

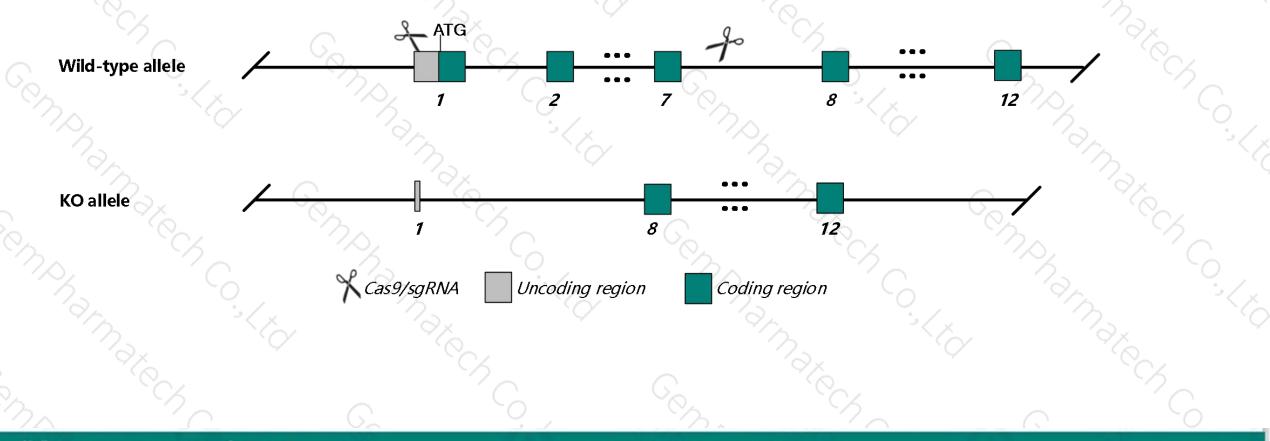






400-9660890

This model will use CRISPR/Cas9 technology to edit the Shc1 gene. The schematic diagram is as follows:



江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.



➤ The Shc1 gene has 10 transcripts. According to the structure of Shc1 gene, exon1-exon7 of Shc1-202(ENSMUST00000094378.9) 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 *Shc1* gene. The brief process is as follows: CRISPR/Cas9 system 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.



> According to the existing MGI data, homozygotes with a targeted mutation of the exon encoding the CH2 region show an extended life span, reduced cellular sensitivity to oxidative stress and UV irradiation, and resistance to diet-induced atherogenesis. Homozygotes lacking all three isoforms die around E11.5 with cardiovascular defects. The distance between *Cks1b* and *Shc1-202* is about 3kb, and the 5-terminal regulation of *Cks1b* may be affect. > The Shc1 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.

## Gene information (NCBI)



Shc1 src homology 2 domain-containing transforming protein C1 [ Mus musculus (house mouse) ]

Gene ID: 20416, updated on 6-Sep-2020

### Summary

Official SymbolShc1 provided by MGIOfficial Full Namesrc homology 2 domain-containing transforming protein C1 provided by MGIPrimary sourceMGI:MGI:98296See relatedEnsembl:ENSMUSG00000042626Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae;<br/>Murinae; Musi NusAlso knownasSh; p6; Shc, p66; ShcA; p66s, p66shcExpressionUbiquitous expression in subcutaneous fat pad adult (RPKM 33.0), ovary adult (RPKM 32.1) and 28 other tissues See more<br/>human all

### 江苏集萃药康生物科技股份有限公司

### GemPharmatech Co., Ltd.

### 400-9660890

(2) ?

## **Transcript information (Ensembl)**



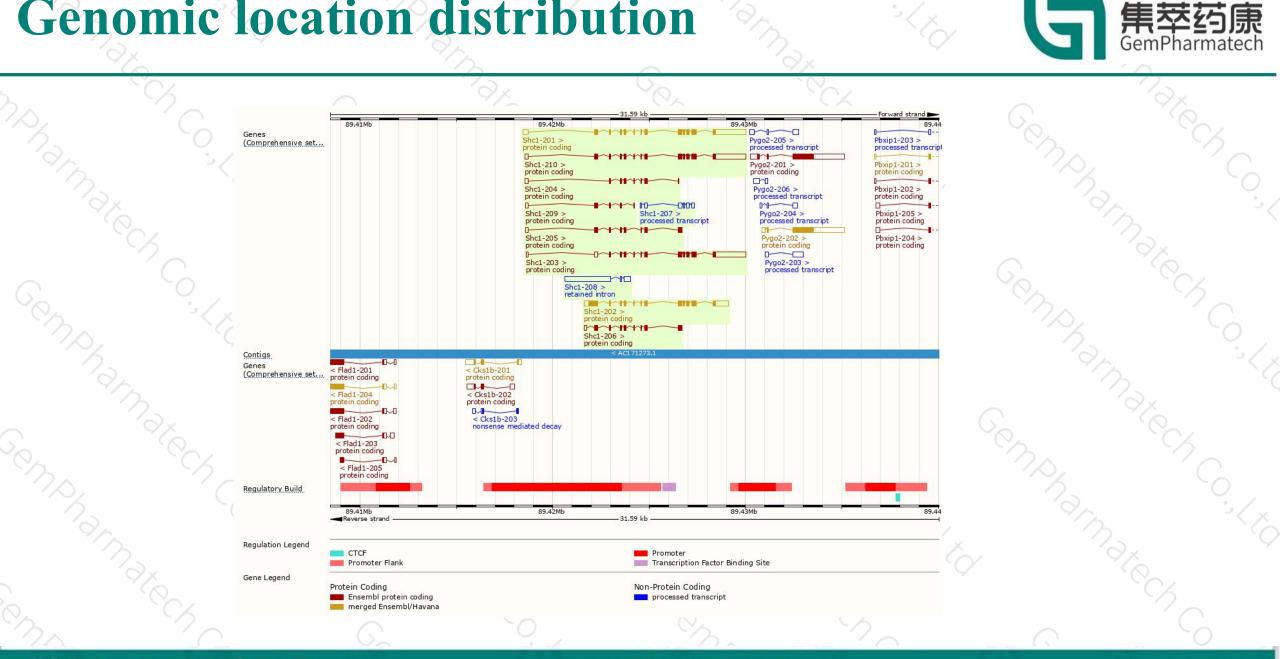
## The gene has 10 transcripts, all transcripts are shown below:

Name 🖕	Transcript ID	bp 🛊	Protein A	Biotype	CCDS 🔶	UniProt 🛊	Flags
Shc1-209	ENSMUST00000154791.7	497	<u>118aa</u>	Protein coding		<u>D3YXZ9</u> ₽	CDS 3' incomplete TSL:3
Shc1-204	ENSMUST00000125036.7	679	<u>172aa</u>	Protein coding	5. <del>4</del> 0	<u>D3YZV5</u> ₽	CDS 3' incomplete TSL:3
Shc1-205	ENSMUST00000128238.7	906	<u>262aa</u>	Protein coding	( <del>•</del> )	D3Z218	CDS 3' incomplete TSL:3
Shc1-206	ENSMUST00000137793.1	951	<u>280aa</u>	Protein coding	23 <b>4</b> 2	D3Z5U6@	CDS 3' incomplete TSL:3
Shc1-203	ENSMUST00000107417.8	3079	<u>424aa</u>	Protein coding	1	P98083	TSL:1 GENCODE basic
Shc1-201	ENSMUST0000039110.11	3250	<u>469aa</u>	Protein coding	<u>CCDS17508</u> 교	P98083	TSL:1 GENCODE basic APPRIS P3
Shc1-210	ENSMUST00000191485.6	3154	<u>469aa</u>	Protein coding	CCDS17508	P98083	TSL:1 GENCODE basic APPRIS P3
Shc1-202	ENSMUST0000094378.9	2617	<u>579aa</u>	Protein coding	<u>CCDS50962</u> 교	P98083	TSL:1 GENCODE basic APPRIS ALT1
Shc1-207	ENSMUST00000146306.1	00146306.1 748		Processed transcript		7	TSL:3
Shc1-208	ENSMUST00000153334.1	2775	No protein	Retained intron	(1 <del>9</del> )	*	TSL:1

The strategy is based on the design of *Shc1-202* transcript, the transcription is shown below:



## **Genomic location distribution**



江苏集萃药康牛物科技股份 限公司

### GemPharmatech Co., Ltd.

## **Protein domain**



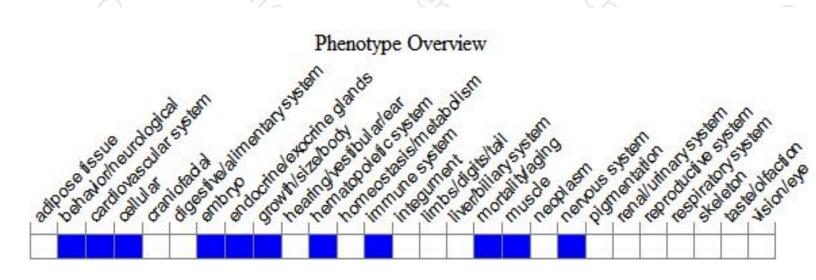
5.	19		2 CK	17 SZ	10	Gen a	°°	
ç	ENSMUSP00000091 MobiDB lite Low complexity (Seg) Superfamily		SSF50729			SH2 domain superfamily		
	<u>SMART</u>		PTB/PI domain			SH2 domain	×	
	Prints Pfam.		Phosphotyrosine interaction PTB/PI domain	n domain, Shc-like		SH2 domain SH2 domain	° ~	
` '	PROSITE profiles		PTB/PI domain			SH2 domain	- ~	
	PANTHER	PTHR10337						
	Gene3D	SHC-transforming protein 1	PH-like domain superfamily			SH2 domain superfamily		
	CDD.		Phosphotyrosine interaction d	omain, Shc-like		SH2 adaptor protein C, SH2	domain 6	
	All sequence SNPs/i	Sequence variants (dbSNP and	d all other sources)	a a	11	1, 1, 11		
	Variant Legend	missense variant		synonymo	us variant			
	Scale bar	0 60	120 180	240 300	360 420	480	579	

江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.

## 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 with a targeted mutation of the exon encoding the CH2 region show an extended life span, reduced cellular sensitivity to oxidative stress and UV irradiation, and resistance to diet-induced atherogenesis. Homozygotes lacking all three isoforms die around E11.5 with cardiovascular defects.

## 江苏集萃药康生物科技股份有限公司

### GemPharmatech Co., Ltd.



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



