

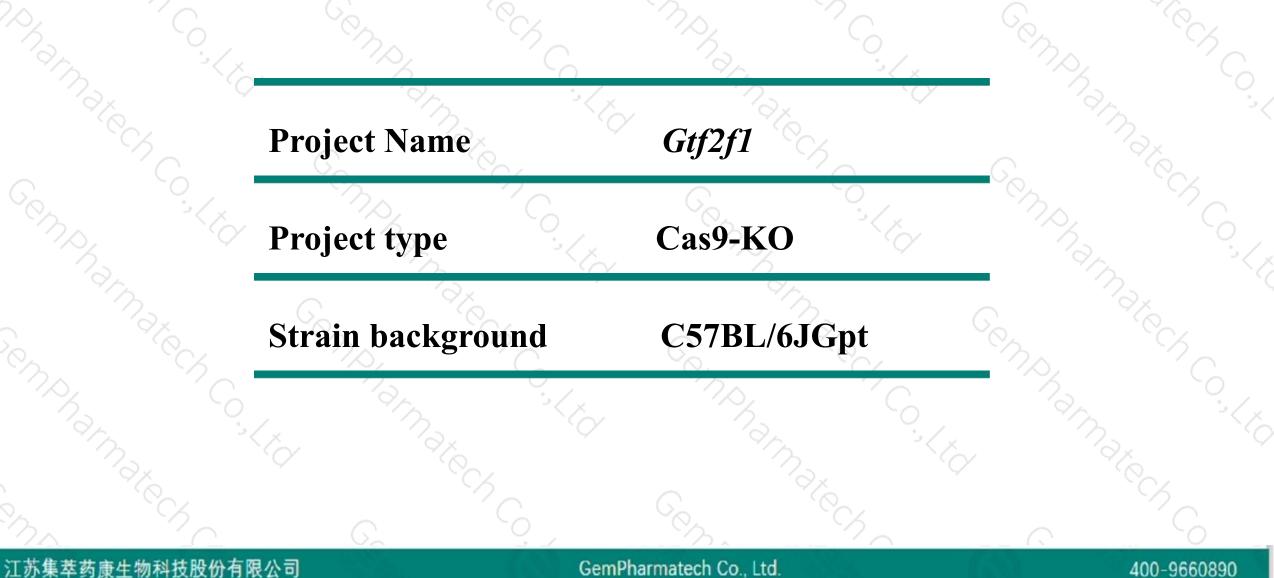
Gtf2f1 Cas9-KO Strategy

Designer: Reviewer. Design Date: Ruirui Zhang Huimin Su

2019-8-23

Project Overview



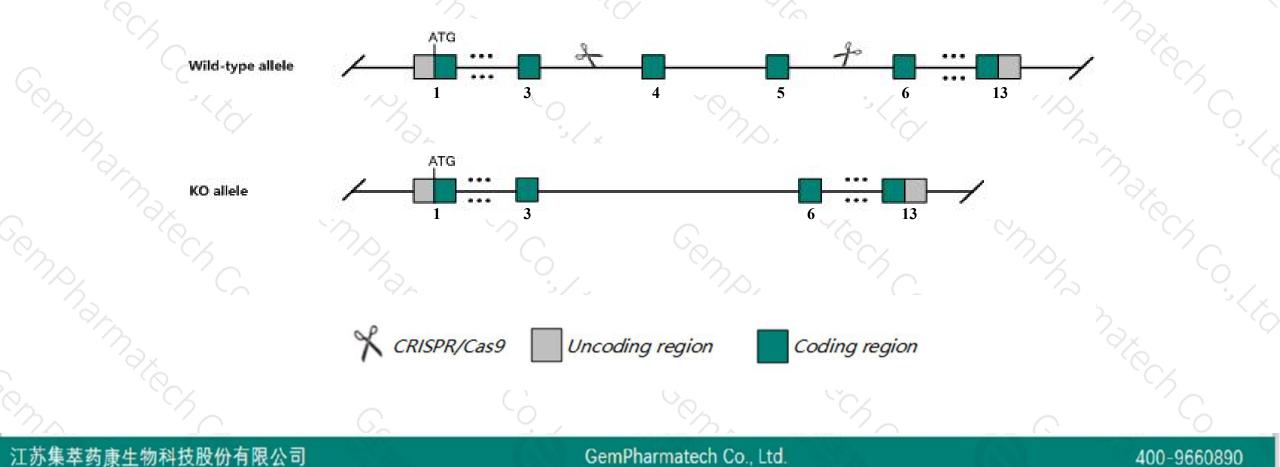


GemPharmatech Co., Ltd.

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gtf2f1* gene. The schematic diagram is as follows:





- The *Gtf2f1* gene has 1 transcript. According to the structure of *Gtf2f1* gene, exon4-exon5 of *Gtf2f1-201* (ENSMUST0000002733.6) transcript is recommended as the knockout region. The region contains 365bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Gtf2f1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Gtf2f1* gene is located on the Chr17. 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.
- The distance between exon4 of *Gtf2f1* gene and *Gm17949* is about 3.5kb, this strategy may affect the 5 terminals regulatory function of *Gm17949* gene.
- 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)



☆ ?

Gtf2f1 general transcription factor IIF, polypeptide 1 [Mus musculus (house mouse)]

Gene ID: 98053, updated on 12-Aug-2019

Summary

Official Symbol Gtf2f1 provided by MGI Official Full Name general transcription factor IIF, polypeptide 1 provided by MGI Primary source MGI:MGI:1923848 Ensembl:ENSMUSG0000002658 See related Gene type protein coding RefSeg status PROVISIONAL Organism Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Lineage Murinae; Mus; Mus Also known as RAP74; C76800; 2810405L04Rik Expression Ubiquitous expression in testis adult (RPKM 78.6), liver E14 (RPKM 38.3) and 28 other tissues See more Orthologs human all

江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.

400-9660890

Transcript information (Ensembl)

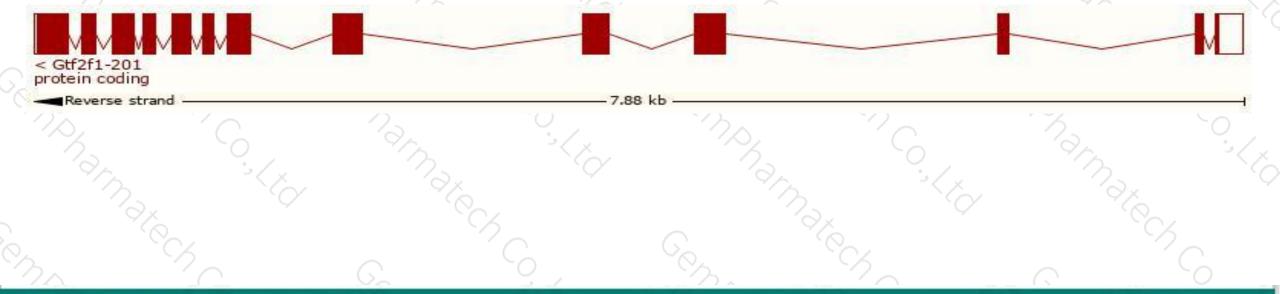


400-9660890

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Gtf2f1-201	ENSMUST0000002733.6	1717	<u>508aa</u>	Protein coding	CCDS37667	Q3THK3	TSL:1 GENCODE basic APPRIS P1	E

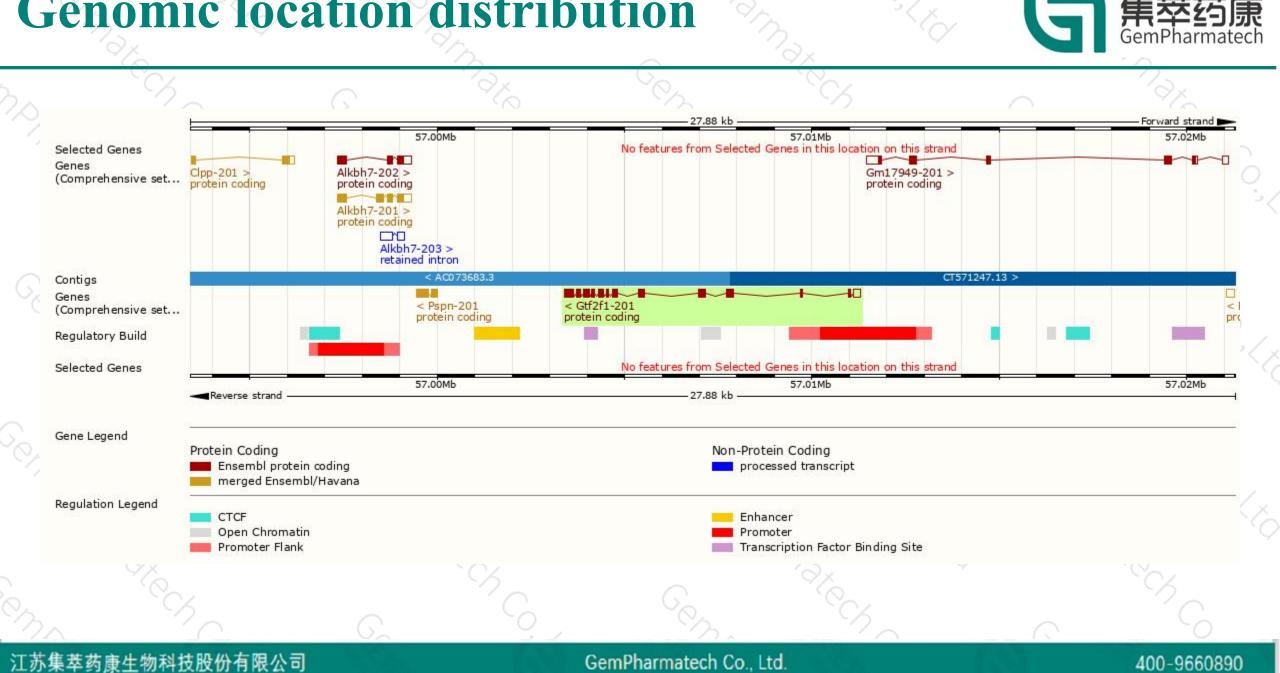
The strategy is based on the design of *Gtf2f1-201* transcript, The transcription is shown below



江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.

Genomic location distribution



集萃药

Protein domain



		Co Sector	CM SS	с С	n phan		COMP.	°°G C	
	ENSMUSP00000002 MobiDB lite Low complexity (Seg) Superfamily	-	, Rap30/Rap74, interaction				Winged	helix DNA-binding dom	
	Pfam	Transcription initiation fa	ictor IIF, alpha subunit						
	PANTHER	PTHR13011:SF0							
26	Gene3D CDD		factor IIF, alpha subunit				Winged hel	ix-like DNA-binding dom	
		cd00240	CND and all athen assured					2/	
	All sequence SNPs/i	Sequence variants (db:	SNP and all other sources)	C C (D)	L. L. L.	1	I.		
	Variant Legend	missense variant			synoi	nymous variant	ious variant		
	Scale bar		50 120	180	240	300 360	420	508	
	Charmar	NO. KA	harmate		C'NDA	And And	y harma		
	^°Ç	×	G_	6	Con .		0	30	

江济



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



