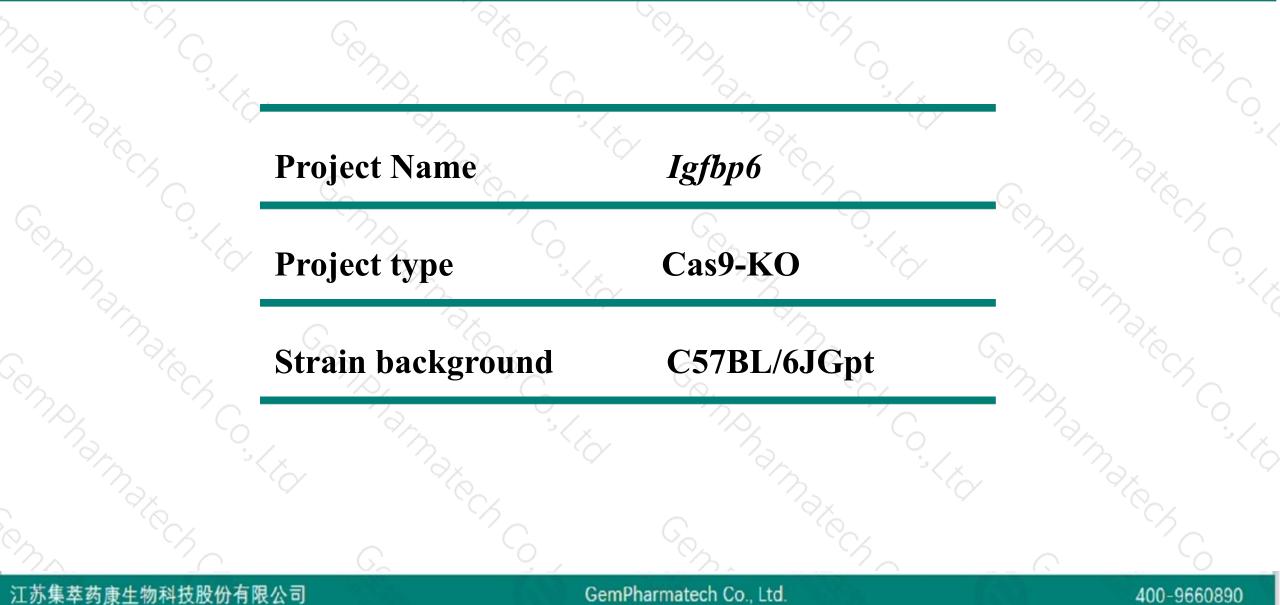


Igfbp6 Cas9-KO Strategy

Designer: Reviewer: Design Date: Yang Zeng Jing Jin 2019-10-31

Project Overview

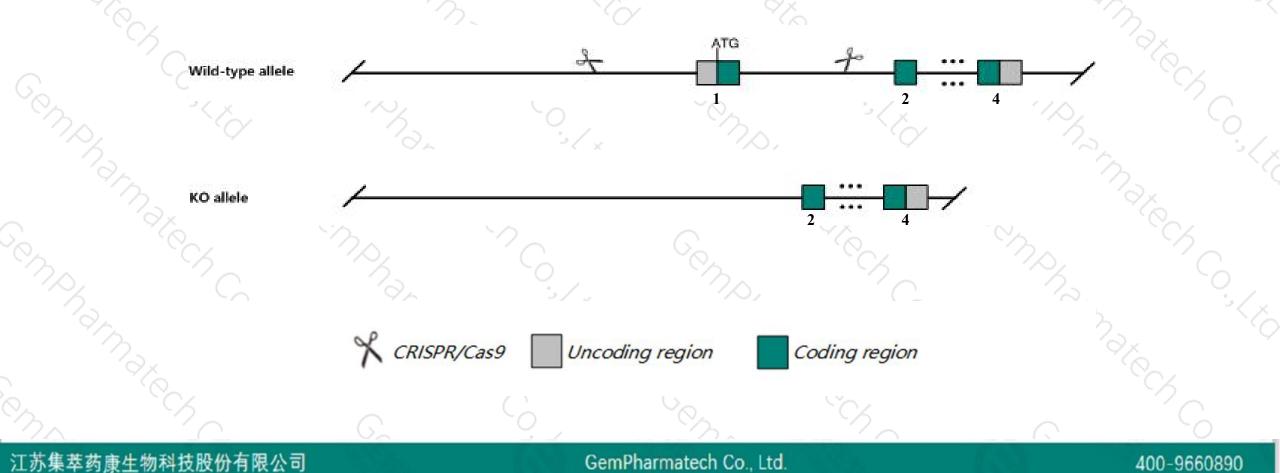




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Igfbp6* gene. The schematic diagram is as follows:





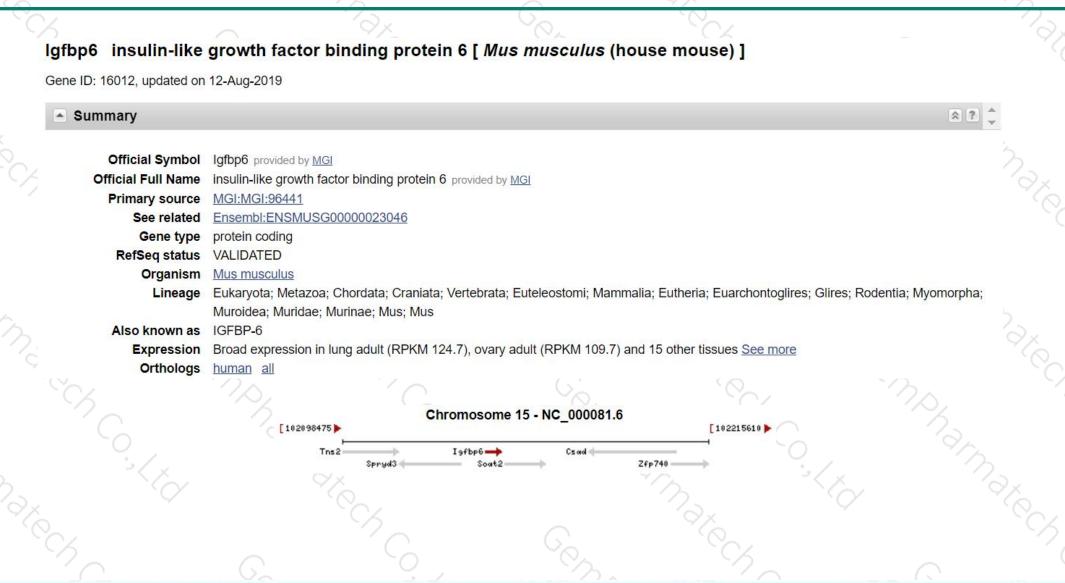
- The Igfbp6 gene has 1 transcript. According to the structure of Igfbp6 gene, exon1 of Igfbp6-201 (ENSMUST00000023807.6) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Igfbp6 gene. The brief process is as follows: CRISPR/Cas9 system

- The *Igfbp6* gene is located on the Chr15. 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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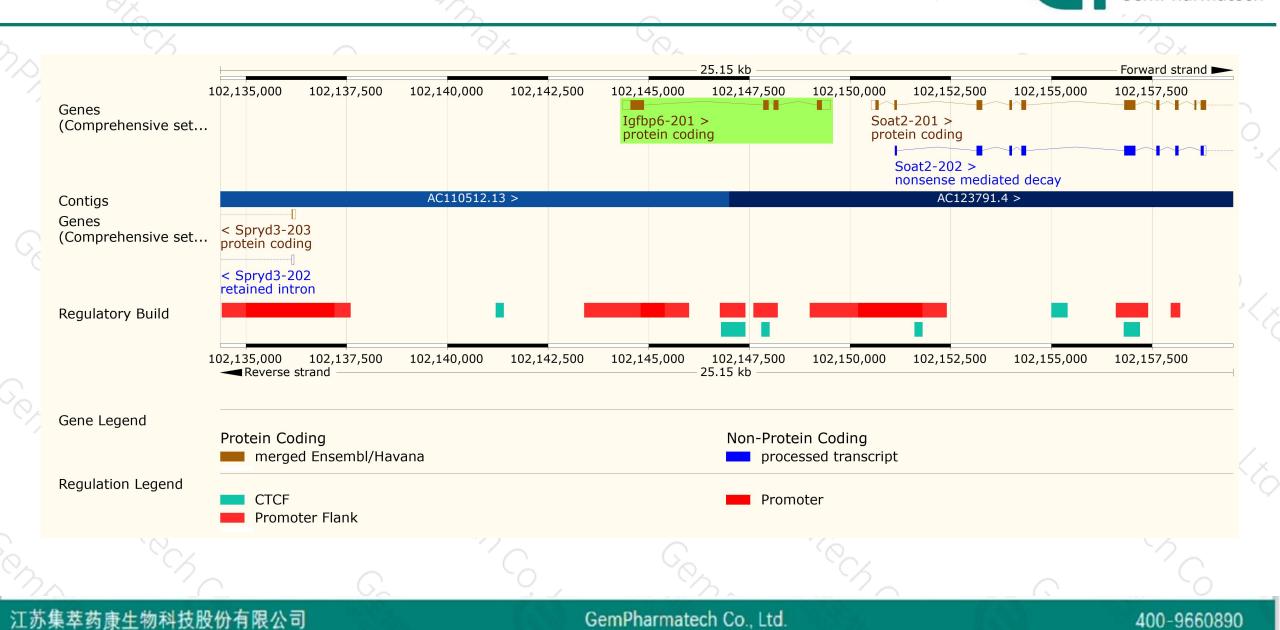
The gene has 1 transcript, and the transcript is shown below:

Name 🖕	Transcript ID 💧	bp 🍦	Protein 🖕	Translation ID	Biotype 🍦	CCDS	UniProt 🖕		Flags	4
lgfbp6-201	ENSMUST0000023807.6	1112	<u>238aa</u>	ENSMUSP0000023807.6	Protein coding	<u>CCDS27871</u> &	<u>P47880</u>	TSL:1	GENCODE basic	APPRIS P1

The strategy is based on the design of Igfbp6-201 transcript, The transcription is shown below



Genomic location distribution



Protein domain



	Transmembrane heli MobiDB lite Low complexity (Seg)		_					_					
	Cleavage site (Sign Superfamily			Growth fac	ctor receptor cystein	e-rich domain superfamily	т	hyroglobulin type-	1 superfamily				
	SMART	Ins	ulin-like growth factor-bi	nding protein, IGFBP)				Thyrog	globulin type-1			
	Prints			Insulin-		inding protein family 1-6, chord					<u> </u>		
	Pfam				Insulin-like	e growth factor-binding protein	6						
	PROSITE profiles	Total						Thyroglobu				_	
	PROSITE promes	Insui	n-like growth factor-bind	ing protein, IGFBP				Thyroglobulin	type-1				
1	PROSITE patterns			Insulin-like growth	factor binding protei	in, N-terminal, Cys-rich conser	ved site		Thyrog	lobulin type-1) 's /
	PANTHER	Insulin-like growth factor-bind	ing protein 6										
		Insulin-like growth factor-bind Insulin-like growth factor bind											
	PANTHER Gene3D			4.10.40.20			Thy	roglobulin type-1	superfamily				
				4.10.40.20			Thy	roglobulin type-1 Thyroglobuli					
	Gene3D		ing protein	4.10.40.20			Thy					- 2	
	Gene3D CDD	Insulin-like growth factor bind	ing protein	4.10.40.20		ous variant			n type-1	1			
	Gene3D CDD All sequence SNPs/i	Insulin-like growth factor bind Sequence variants (dbSNP	ing protein	4.10.40.20	synonymo 80	ous variant			n type-1	200		238	
	Gene3D CDD All sequence SNPs/i Variant Legend	Insulin-like growth factor bind	and all other sources)		synonymo			Thyroglobuli	n type-1	200	• •	238	
	Gene3D CDD All sequence SNPs/i Variant Legend	Insulin-like growth factor bind	and all other sources)		synonymo			Thyroglobuli	n type-1	200		238	
	Gene3D CDD All sequence SNPs/i	Insulin-like growth factor bind Sequence variants (dbSNP	ing protein	4.10.40.20		ous variant			n type-1	1	1	_	20**

江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.

400-9660890



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



