

# Klf4 Cas9-KO Strategy

Designer: Reviewer:

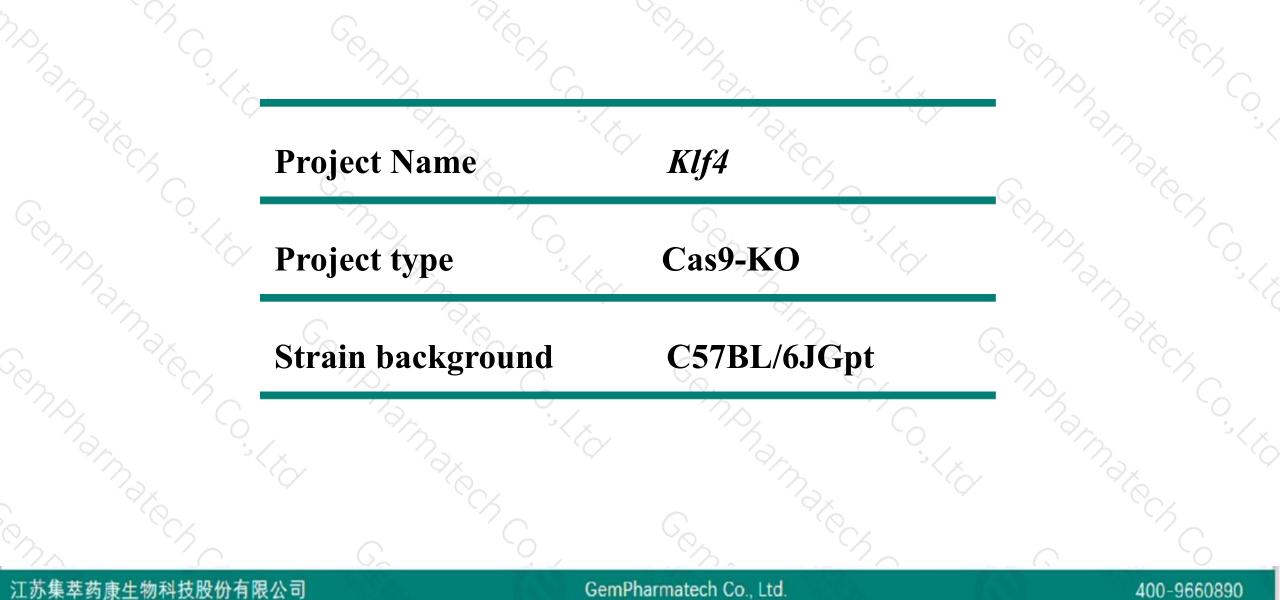
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Design Date:

Huan Wang Huan Fan 2020-3-9

### **Project Overview**

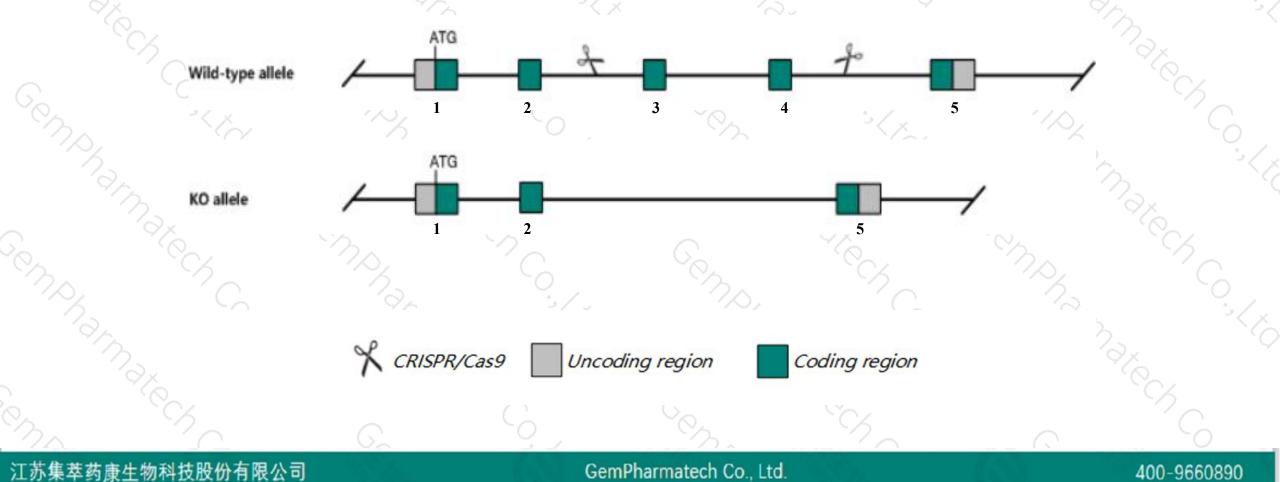




### **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the *Klf4* gene. The schematic diagram is as follows:





- The Klf4 gene has 3 transcripts. According to the structure of Klf4 gene, exon3-exon4 of Klf4-201 (ENSMUST00000107619.2) transcript is recommended as the knockout region. The region contains 1150bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Klf4 gene. The brief process is as follows: CRISPR/Cas9 system w

- According to the existing MGI data, Homozygotes for targeted null mutations die shortly after birth due to a skin defect that results in loss of fluids. Mutants also show a dramatic decrease in the number of goblet cells of the colon.
- The *Klf4* gene is located on the Chr4. 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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#### KIf4 Kruppel-like factor 4 (gut) [Mus musculus (house mouse)]

Gene ID: 16600, updated on 26-Mar-2019

#### Summary

Official Symbol	KIf4 provided by MGI
Official Full Name	Kruppel-like factor 4 (gut) provided by MGI
Primary source	MGI:MGI:1342287
See related	Ensembl:ENSMUSG0000003032
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	EZF, Gklf, Zie
Expression	Biased expression in colon adult (RPKM 221.0), stomach adult (RPKM 155.0) and 9 other tissues See more
Orthologs	human all

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## **Transcript information (Ensembl)**



#### The gene has 3 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
KIf4-201	ENSMUST00000107619.2	3029	<u>483aa</u>	Protein coding	CCDS18195	F2YID5 Q60793	TSL:1 GENCODE basic APPRIS P1
KIf4-203	ENSMUST00000132746.1	859	<u>82aa</u>	Protein coding		B7ZCH1	CDS 3' incomplete TSL:3
KIf4-202	ENSMUST00000129250.1	522	<u>160aa</u>	Protein coding	620	B7ZCH2	CDS 3' incomplete TSL:2

The strategy is based on the design of *Klf4-201* transcript, The transcription is shown below

#### < Klf4-201 protein coding

Reverse strand

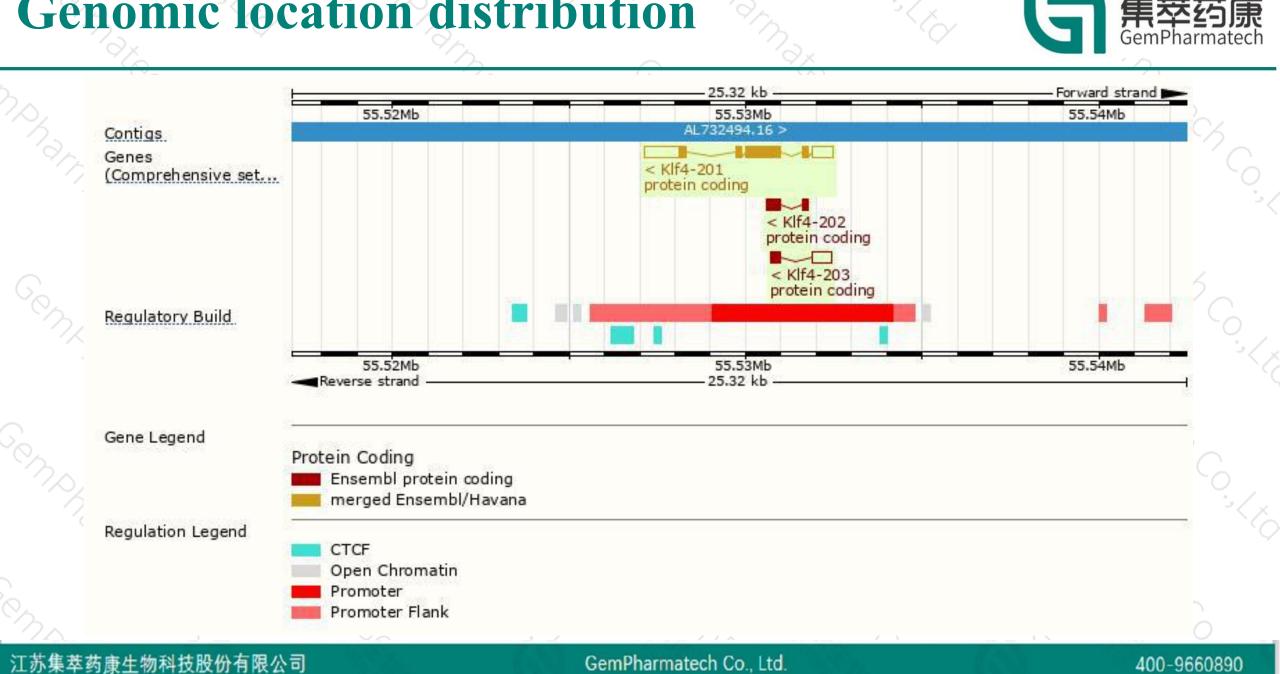
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5.32 kb

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### **Genomic location distribution**



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### **Protein domain**



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	Low complexity (Seq)			Sec.						
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	Pfam.							Zinc finger C2		
	PROSITE profiles							Zinc finger C2	H2-type	
	PROSITE patterns							Zinc finger C2	H2-type	·. /
	PANTHER	PTHR232351SF52	2							
		PTHR23235								
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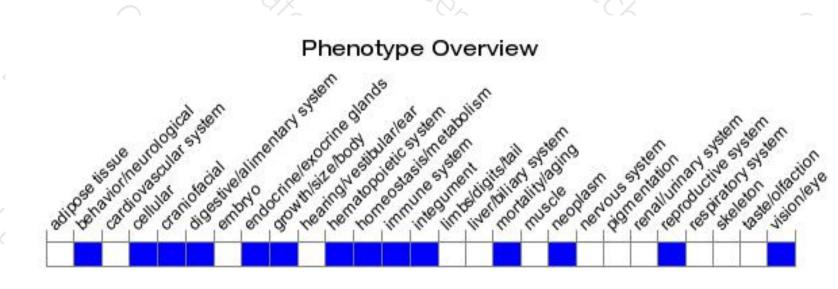
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### Mouse phenotype description(MGI)





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 targeted null mutations die shortly after birth due to a skin defect that results in loss of fluids. Mutants also show a dramatic decrease in the number of goblet cells of the colon.



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



