

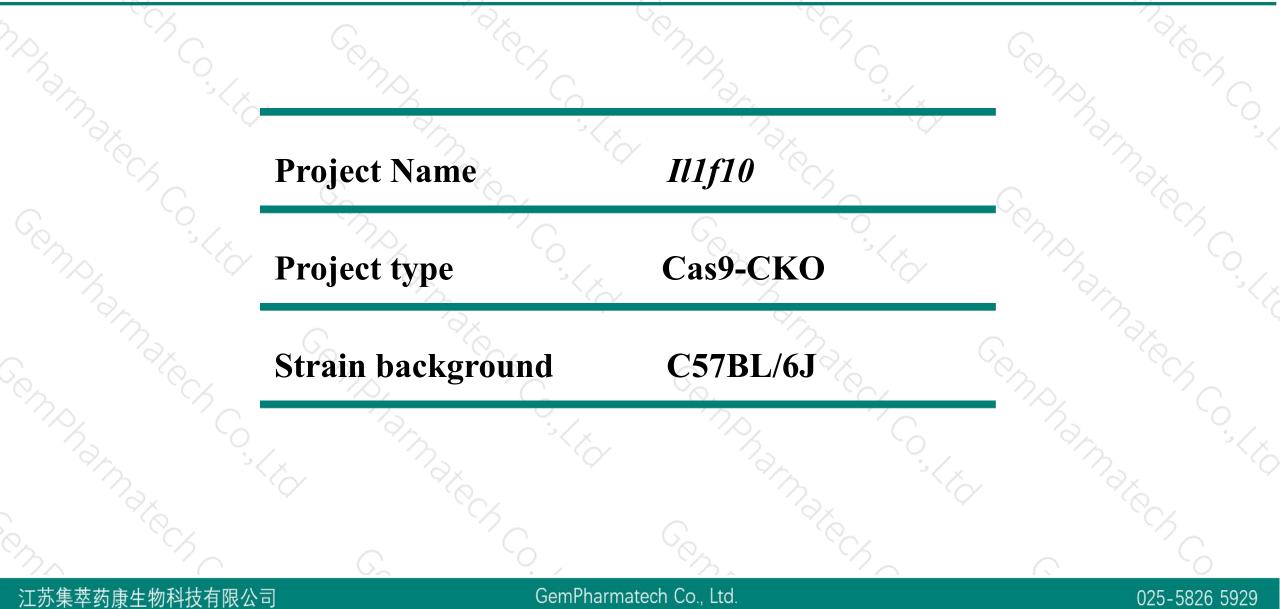
IIIf10 Cas9-CKO Strategy Romphamater Control

Enphamatech C. Lt. JiaYu

empharmatech Co

Project Overview



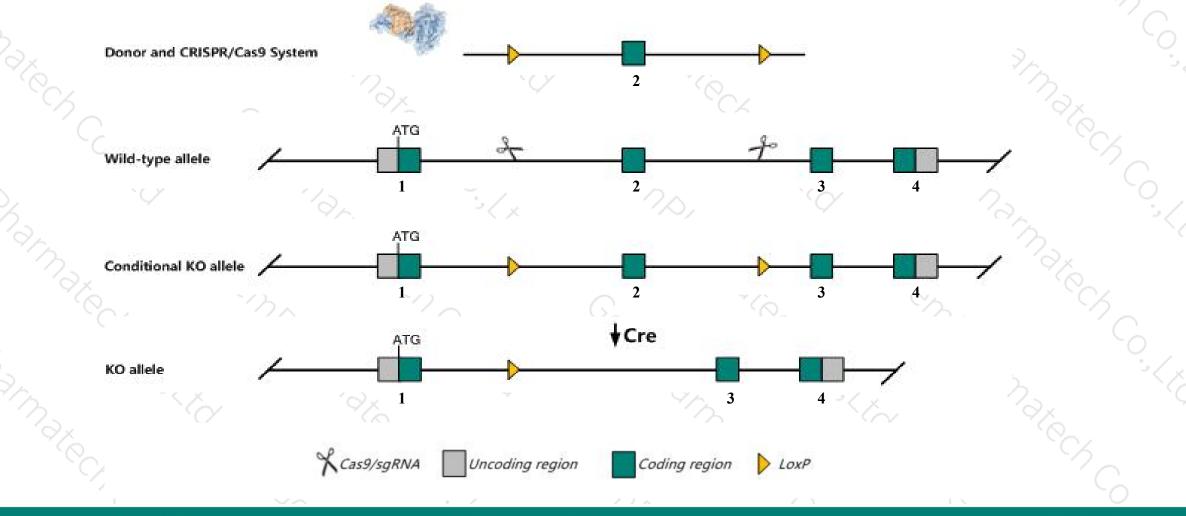


Conditional Knockout strategy



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This model will use CRISPR/Cas9 technology to edit the *Illf10* gene. The schematic diagram is as follows:



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The *Illf10* gene has 1 transcript. According to the structure of *Illf10* gene, exon2 of *Illf10-201* (ENSMUST00000058056.1) transcript is recommended as the knockout region. The region contains 86bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Illf10* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA and Donor were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

The flox mice was 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.



- The *Illf10* 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)



☆ ?

II1f10 interleukin 1 family, member 10 [Mus musculus (house mouse)]

Gene ID: 215274, updated on 31-Jan-2019

Summary

Official Symbol	II1f10 provided by MGI
Official Full Name	interleukin 1 family, member 10 provided by MGI
Primary source	MGI:MGI:2652548
See related	Ensembl:ENSMUSG0000046845
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
	Muroidea; Muridae; Murinae; Mus; Mus
Expression	Low expression observed in reference datasetSee more
Orthologs	human all

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

Name	ne Transcript ID		Protein	Biotype Protein coding	CCDS	UniProt	Flags	
1f10-201	ENSMUST0000058056.1	538 <u>152aa</u>			CCDS15735	Q3KNF2 Q8R459	TSL:1 GENCODE basic APPRIS P1	
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e strateg	y is based on the design	of <i>111</i>	<i>f10-201</i> 1	transcript,The	transcription	is shown below		
<u>_</u>				2.62	kb		Forward strand	

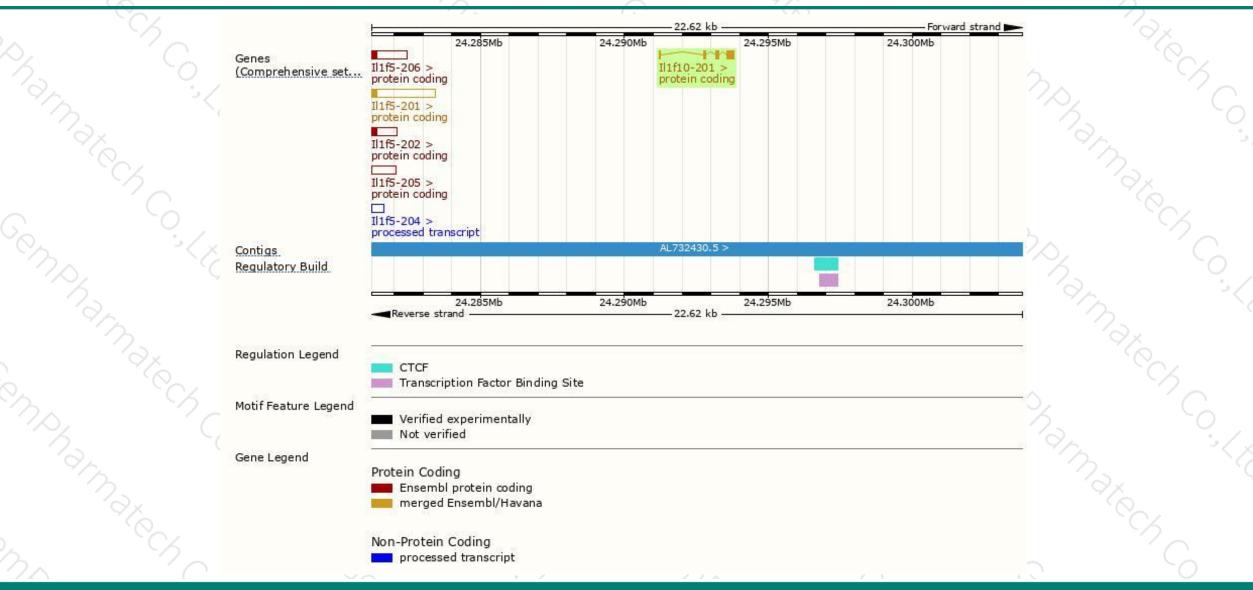
Il1f10-201 > protein coding

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Genomic location distribution



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Protein domain



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ENSMUSP00000063 hmmpanther	Interleukin-1 family	member 10				5	<
Superfamily domains	Cytokine IL1/F	GF					
SMART domains	SM00125						
Prints domain			Interleukin-1 family				8
Pfam domain	Interleukin-1 re	ceptor antagonist/Interleu Interleukin-1				Ň	<->
Gene3D	2.80.10.50		(# C				
All sequence SNPs/i	Sequence variants	(dbSNP and all other s	ources)			1	
Variant Legend	synonymous	variant					0
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If you have any questions, you are welcome to inquire. Tel: 025-5864 1534



