

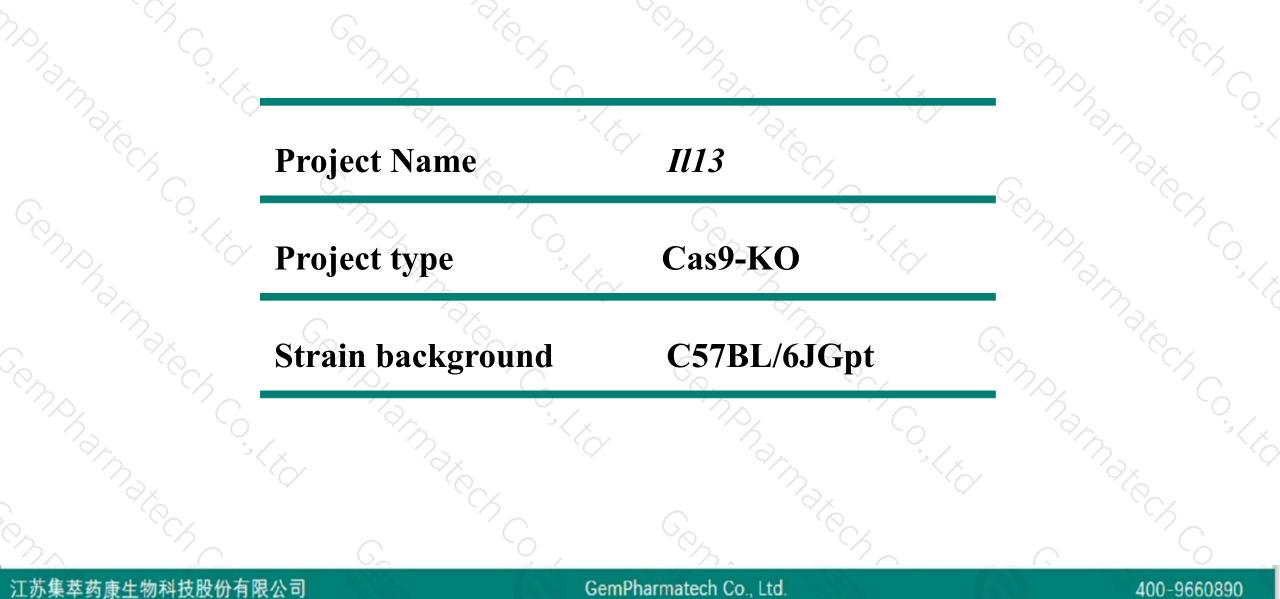
# Il13 Cas9-KO Strategy

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Designer: Xueting Zhang Design Date: 2019-8-3

### **Project Overview**

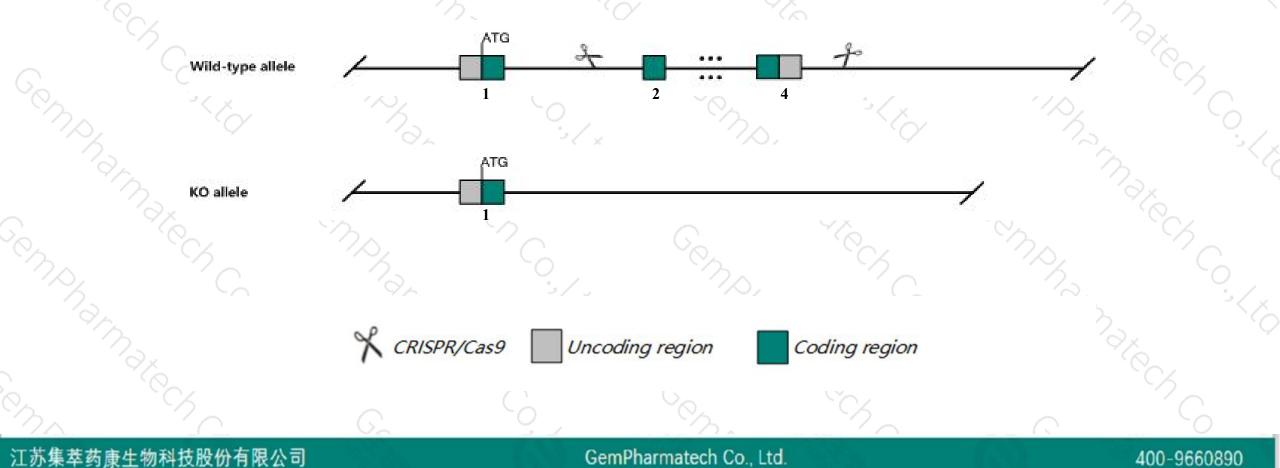




# **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the *Il13* gene. The schematic diagram is as follows:





- The *Il13* gene has 1 transcript. According to the structure of *Il13* gene, exon2-exon4 of *Il13-201* (ENSMUST0000020650.1) transcript is recommended as the knockout region. The region contains 255bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Il13* gene. The brief process is as follows: CRISPR/Cas9 system w

- According to the existing MGI data, Mice homozygous for knock-out alleles exhibit defects in immune system morphology and physiology.
- The *Il13* gene is located on the Chr11. 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)**



☆ ?

### II13 interleukin 13 [Mus musculus (house mouse)]

Gene ID: 16163, updated on 12-Mar-2019

#### Summary

Official SymbolII13 provided by MGIOfficial Full Nameinterleukin 13 provided by MGIPrimary sourceMGI:MGI:96541See relatedEnsembl:ENSMUSG0000020383Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;<br/>Muroidea; Murinae; Mus; MusAlso knownasI-13ExpressionBiased expression in thymus adult (RPKM 2.8), testis adult (RPKM 0.6) and 2 other tissues<br/>Muroidea; Mura adult

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# **Transcript information (Ensembl)**



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

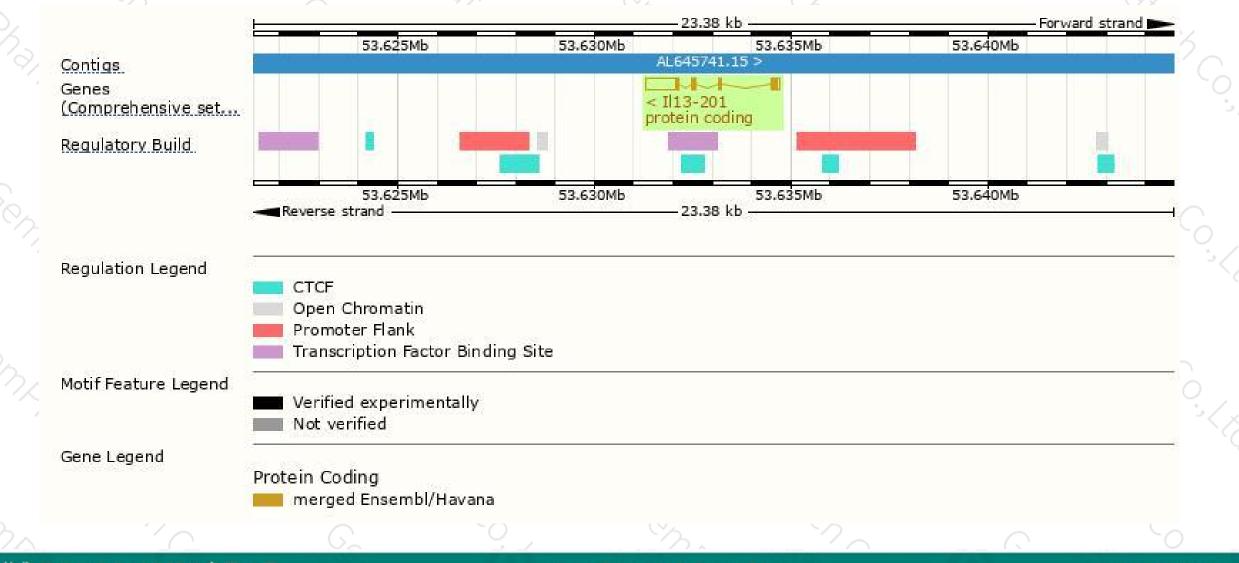
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The strategy is based on the design of <i>II13-201</i> transcript, The transcription is shown below	Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
(113-201 otein coding	13-201	ENSMUST00000020650.1	1205	<u>131aa</u>	Protein coding	CCDS24683	P20109	TSL:1 GENCODE basic APPRIS P1
13-201 tein coding		K. Con		A Ch			K Co	Con Rech
13-201 tein coding						- Chopp		
13-201 tein coding		nate Cont	>			 Sep	ALC CA	Const Conc
	e strate	gy is based on the design	of <i>1113</i> -	201 trans	cript,The transc	cription is show	wn below	
	113-201 tein codi	ng	_					
					3.38 kb			

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### **Genomic location distribution**





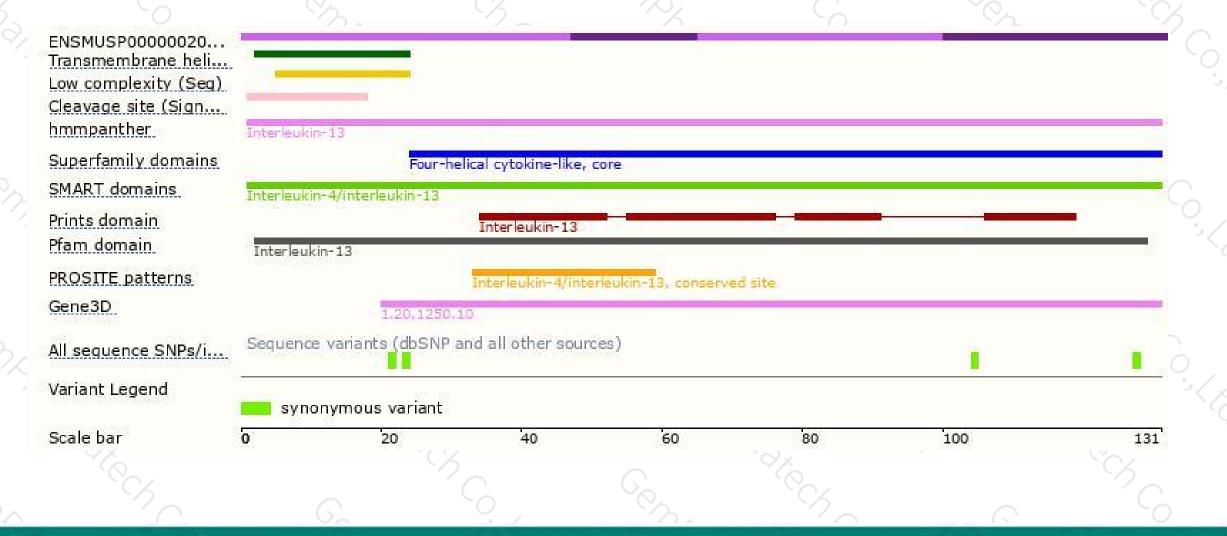
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### **Protein domain**





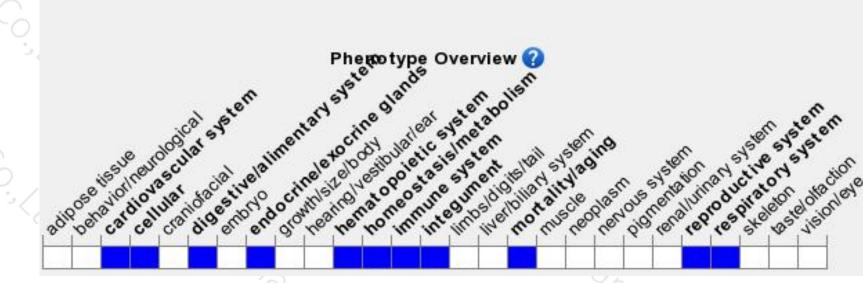
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### Mouse phenotype description(MGI)





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, Mice homozygous for knock-out alleles exhibit defects in immune system morphology and physiology.



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



