

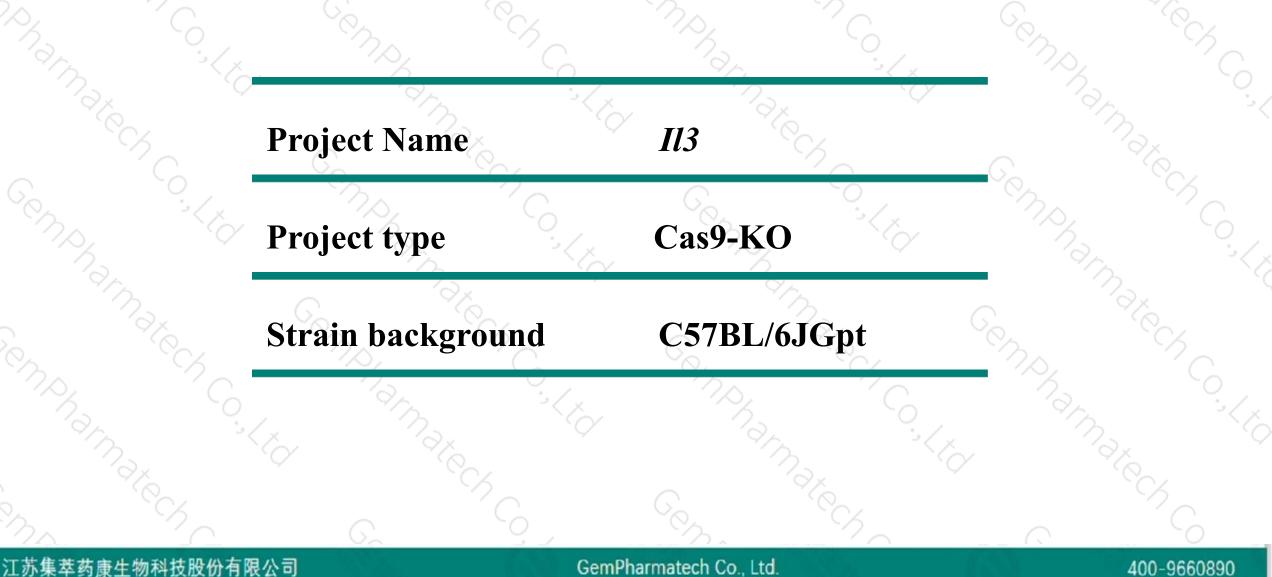
Il3 Cas9-KO Strategy

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Project Overview





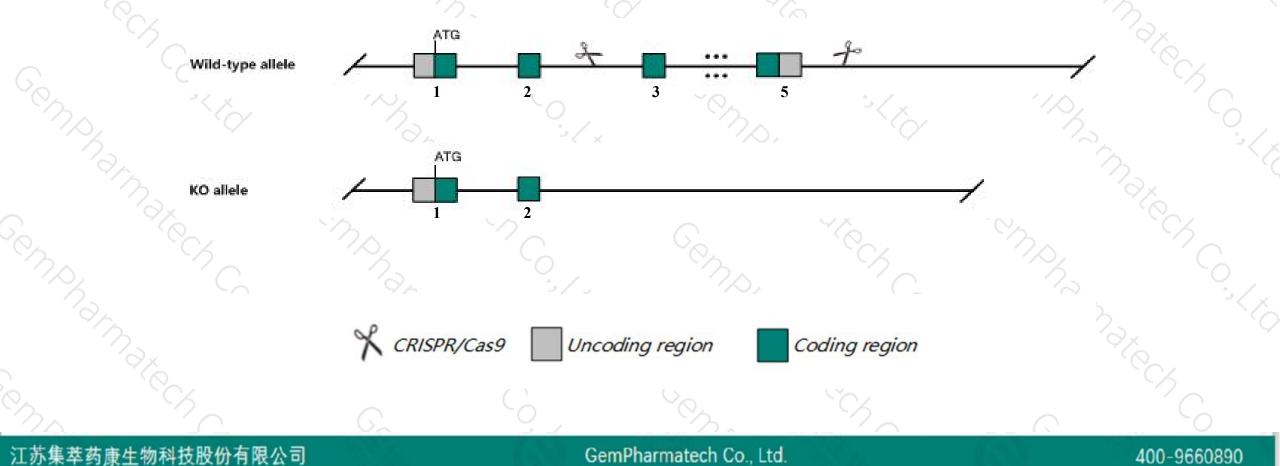
江苏集萃药康生物科技股份有限公司

GemPharmatech Co., Ltd.

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Il3* gene. The schematic diagram is as follows:





- The *Il3* gene has 1 transcript. According to the structure of *Il3* gene, exon3-exon5 of *Il3-201* (ENSMUST00000019058.5) transcript is recommended as the knockout region. The region contains 294bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Il3* gene. The brief process is as follows: CRISPR/Cas9 system we

- According to the existing MGI data, Mice homozygous for disruptions in this gene show abnormal mast cell development. Contact hypersensitivity is reduced and resistance to parasitic invections is increased.
- The *Il3* 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)



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II3 interleukin 3 [Mus musculus (house mouse)]

Gene ID: 16187, updated on 12-Mar-2019

- Summary

Official SymbolII3 provided by MGIOfficial Full Nameinterleukin 3 provided by MGIPrimary sourceMGI:MGI:96552See relatedEnsembl:ENSMUSG00000018914Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
Muroidea; Murinae; Mus; MusAlso knownasBPA, Csfmu, HCGF, II-3, MCGF, PSF

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Transcript information (Ensembl)



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

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Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
13-201	ENSMUST0000019058.5	629	<u>166aa</u>	Protein coding	CCDS24693	P01586 Q5SX77	TSL:1 GENCODE basic APPRIS P1
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		-					
13-201	dian						
Revers	e strand			1.9	8 kb		
		1.2					7.1

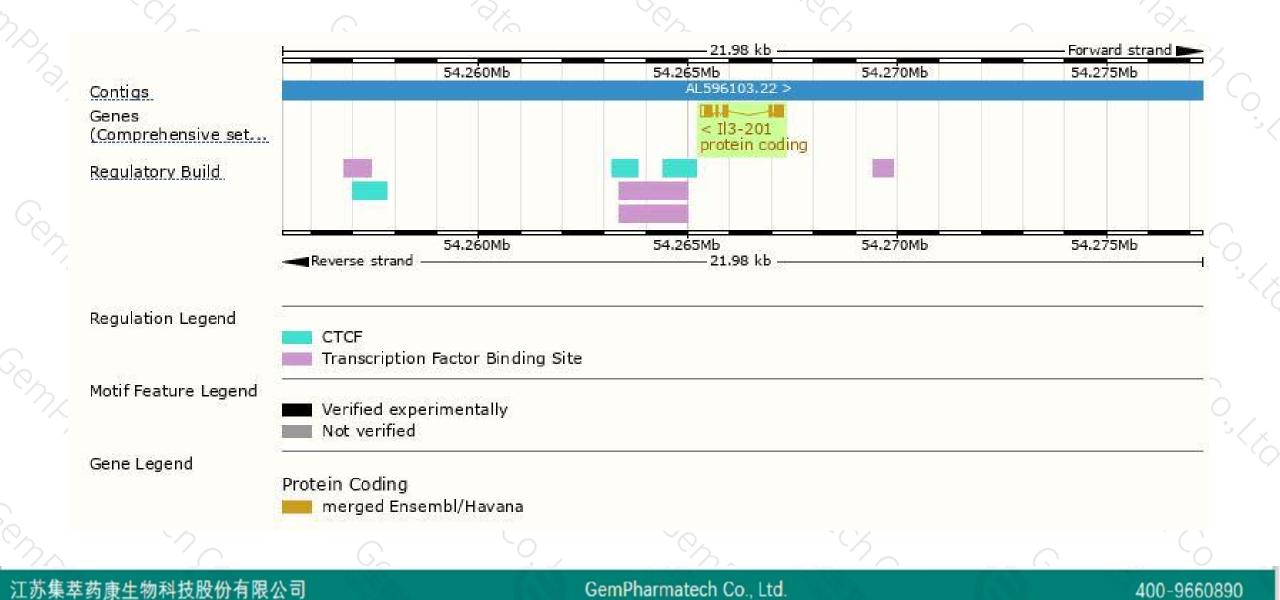
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### **Genomic location distribution**





### **Protein domain**





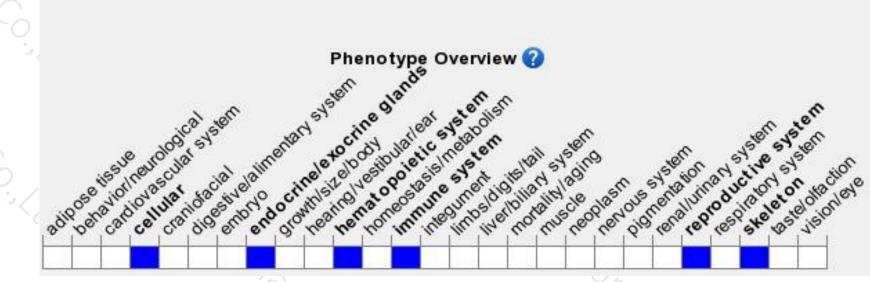
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### Mouse phenotype description(MGI)





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 disruptions in this gene show abnormal mast cell development. Contact hypersensitivity is reduced and resistance to parasitic invections is increased.



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



