

# Il2 Cas9-KO Strategy

Designer: Huan Fan

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# **Project Overview**



**Project Name** 

**Project type** 

Strain background

Cas9-KO

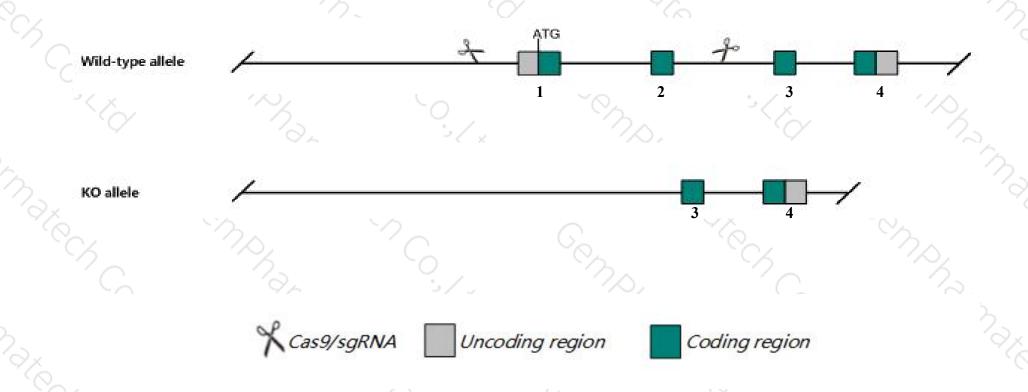
Il2

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Il2* gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The *Il2* gene has 1 transcript. According to the structure of *Il2* gene, exon1-exon2 of *Il2-201*(ENSMUST00000029275.5) 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 *Il2* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6JGpt 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/6JGpt mice.

### **Notice**



- ➤ According to the existing MGI data, Homozygous null mutants develop adult onset autoimmune disease, with 50% mortality by 9 weeks due to hemolytic anemia. Survivors develop inflammatory bowl disease/colitis. Immune system dysregulation and CD4+ T-cell overproduction may be responsible.
- > The *Il2* gene is located on the Chr3. 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.

### Gene information (NCBI)



#### II2 interleukin 2 [Mus musculus (house mouse)]

Gene ID: 16183, updated on 5-Mar-2019

#### Summary

^ ?

Official Symbol II2 provided by MGI

Official Full Name interleukin 2 provided by MGI

Primary source MGI:MGI:96548

See related Ensembl:ENSMUSG00000027720

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 II-2

Expression Low expression observed in reference datasetSee more

Orthologs <u>human all</u>

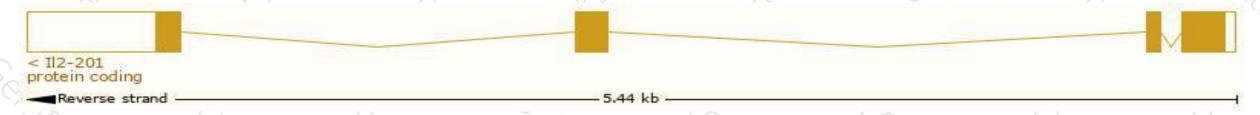
# Transcript information (Ensembl)



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

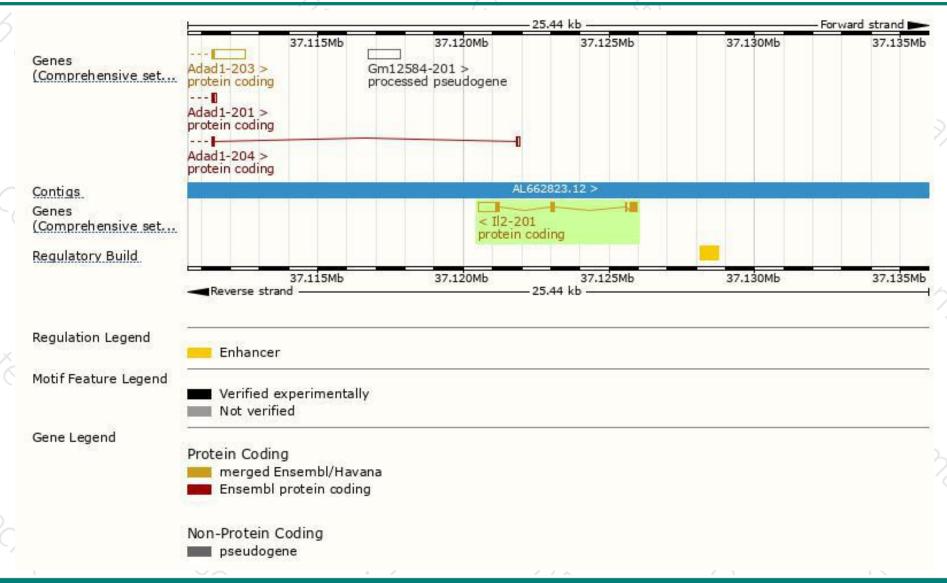
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
112-201	ENSMUST00000029275.5	1144	169aa	Protein coding	CCDS17316	P04351	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Il2-201* transcript, The transcription is shown below



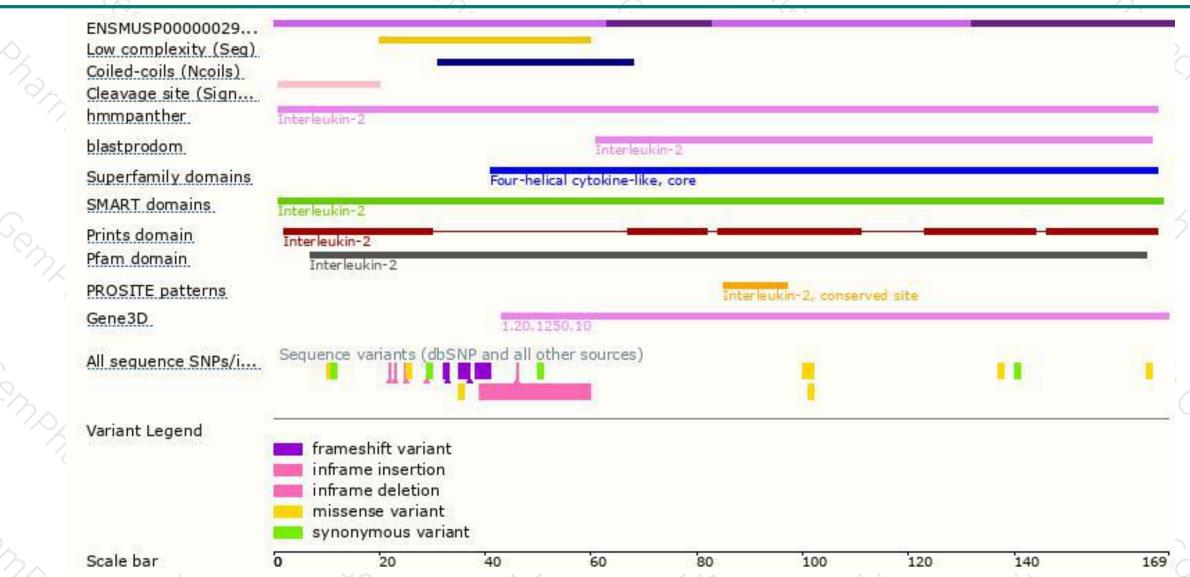
### Genomic location distribution





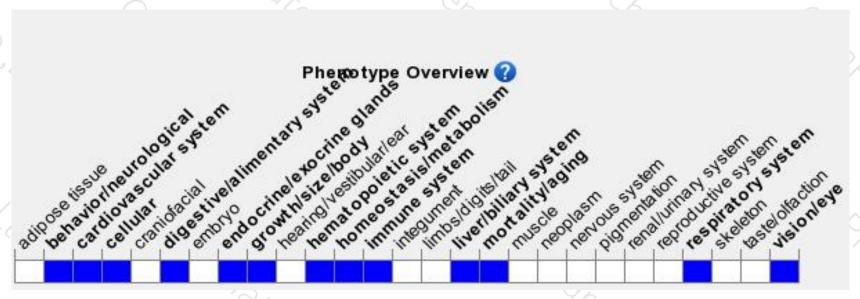
### Protein domain





## 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, Homozygous null mutants develop adult onset autoimmune disease, with 50% mortality by 9 weeks due to hemolytic anemia. Survivors develop inflammatory bowl disease/colitis. Immune system dysregulation and CD4+ T-cell overproduction may be responsible.



If you have any questions, you are welcome to inquire.

Tel: 025-5864 1534





