# Il5 Cas9-CKO Strategy

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**Design Date:** 2019-8-8

## **Project Overview**



**Project Name** 

*Il5* 

**Project type** 

Cas9-CKO

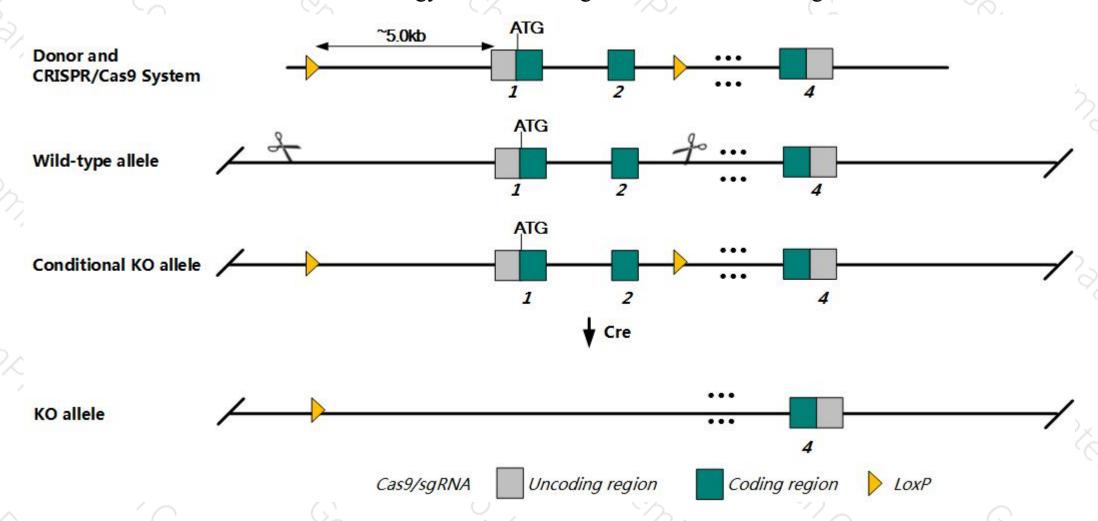
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Il5* gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The *Il5* gene has 1 transcripts. According to the structure of *Il5* gene, the predicted promoter region and exon1-2 of *Il5*-201 (ENSMUST00000048605.2) transcript is recommended as the knockout region. The region contains the predicted promoter sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Il5* gene. The brief process is as follows: gRNA was transcribed in vitro, donor was constructed.Cas9, gRNA and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues or cell types.

### **Notice**



- According to the existing MGI data: Homozygotes for a targeted null mutation exhibit loss of normal airway hyperreactivity resulting from aeroallergen challenge, reduced numbers of CD5+ B cells in the peritoneal cavity at 2 weeks, and some altered responses to schistosomiasis infection.
- ➤ The *Il5* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

# Gene information (NCBI)



#### II5 interleukin 5 [ Mus musculus (house mouse) ]

Gene ID: 16191, updated on 8-Dec-2018

#### Summary

☆ ?

Official Symbol II5 provided by MGI

Official Full Name interleukin 5 provided by MGI

Primary source MGI:MGI:96557

See related Ensembl: ENSMUSG00000036117

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

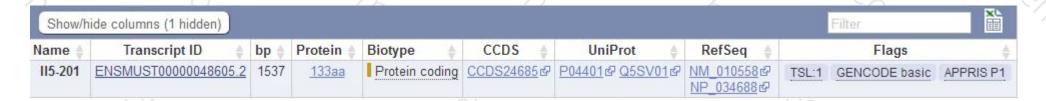
Also known as II-5

Expression Low expression observed in reference dataset See more

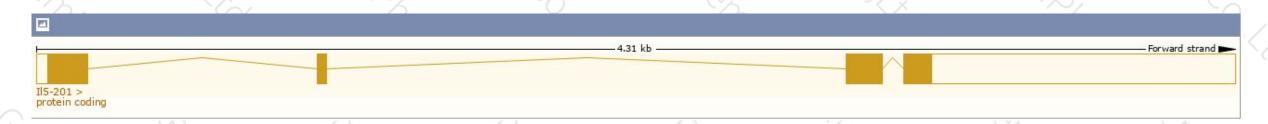
Orthologs human all

# Transcript information (Ensembl 写集萃药康GemPharmatech

The gene has 1 transcripts, and all transcripts are shown below:

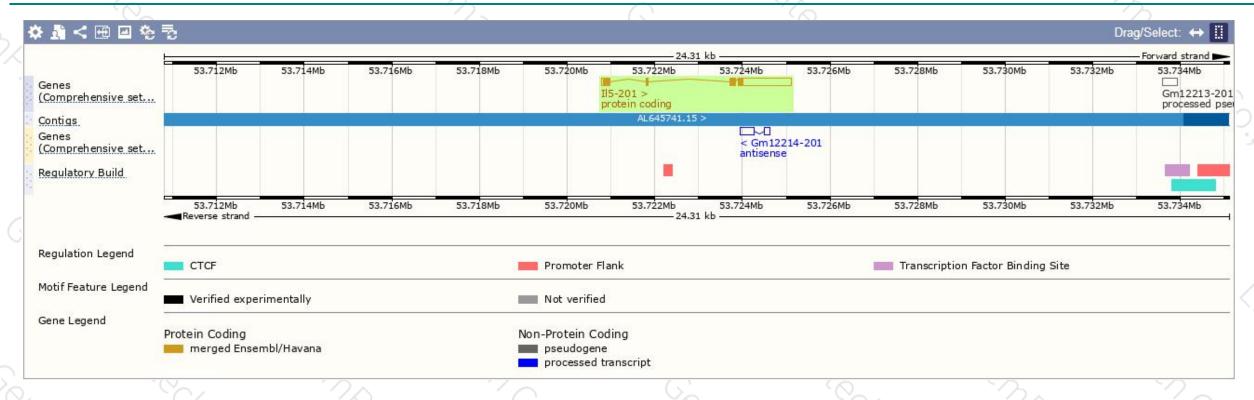


The strategy is based on the design of *Il5-201* transcript, The transcription is shown below



### Genomic location distribution





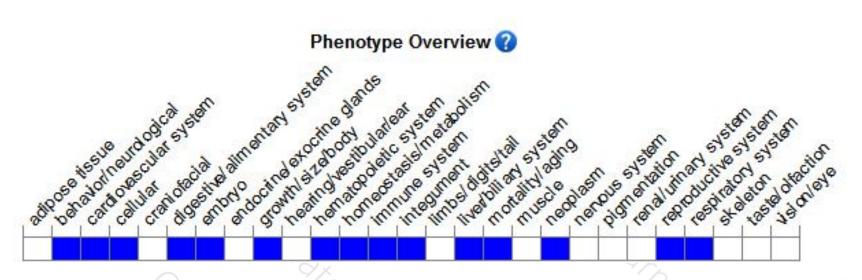
## Protein domain





### Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/) .

Homozygotes for a targeted null mutation exhibit loss of normal airway hyperreactivity resulting from aeroallergen challenge, reduced numbers of CD5+ B cells in the peritoneal cavity at 2 weeks, and some altered responses to schistosomiasis infection.

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





