

# Illf8 Cas9-CKO Strategy

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Design Date: 2019-08-06

# **Project Overview**



**Project Name** 

Il1f8

**Project type** 

Cas9-CKO

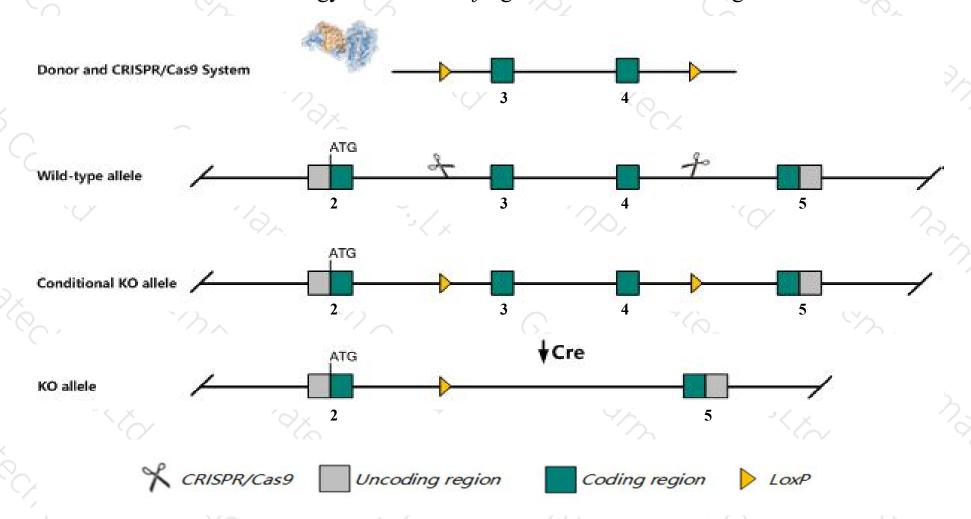
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Il1f*8 gene. The schematic diagram is as follows:



### Technical routes



- ➤ The *Illf8* gene has 1 transcript. According to the structure of *Illf8* gene, exon3-exon4 of *Illf8-201*(ENSMUST00000028363.1) transcript is recommended as the knockout region. The region contains 248bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Il1f8* gene. The brief process is as follows:CRISPR/Cas9 system 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 and cell types.

### **Notice**



- > The *Illf8* 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)



#### II1f8 interleukin 1 family, member 8 [Mus musculus (house mouse)]

Gene ID: 69677, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol II1f8 provided by MGI

Official Full Name interleukin 1 family, member 8 provided by MGI

Primary source MGI:MGI:1916927

See related Ensembl:ENSMUSG00000026985

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

Also known as 2310043N20Rik, II36b

Expression Low expression observed in reference datasetSee more

Orthologs <u>human</u> all

# Transcript information (Ensembl)



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

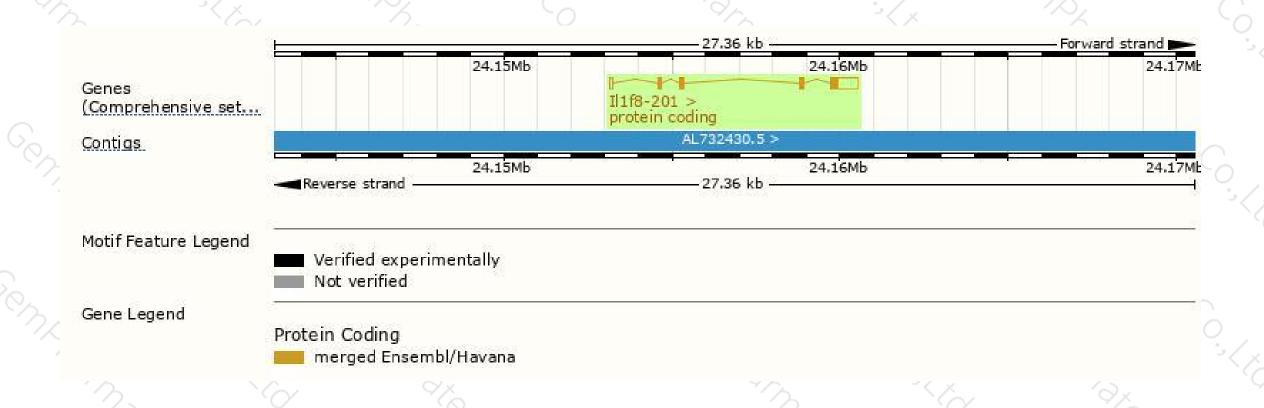
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
II1f8-201	ENSMUST00000028363.1	1217	<u>183aa</u>	Protein coding	CCDS38063	Q08EA6 Q9D6Z6	TSL:1 GENCODE basic APPRIS P1	

The strategy is based on the design of *Illf8-201* transcript, The transcription is shown below



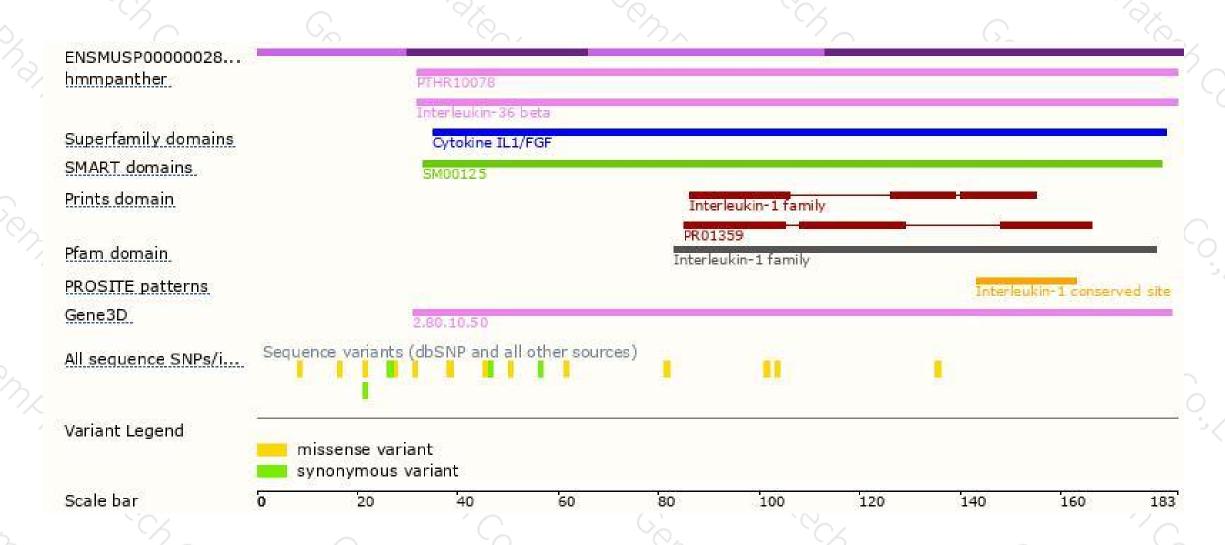
### Genomic location distribution





### Protein domain







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





