

# ***Il17a*** Cas9-KO Strategy

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# Project Overview

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**Project Name**

***Il17a***

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**Project type**

**Cas9-KO**

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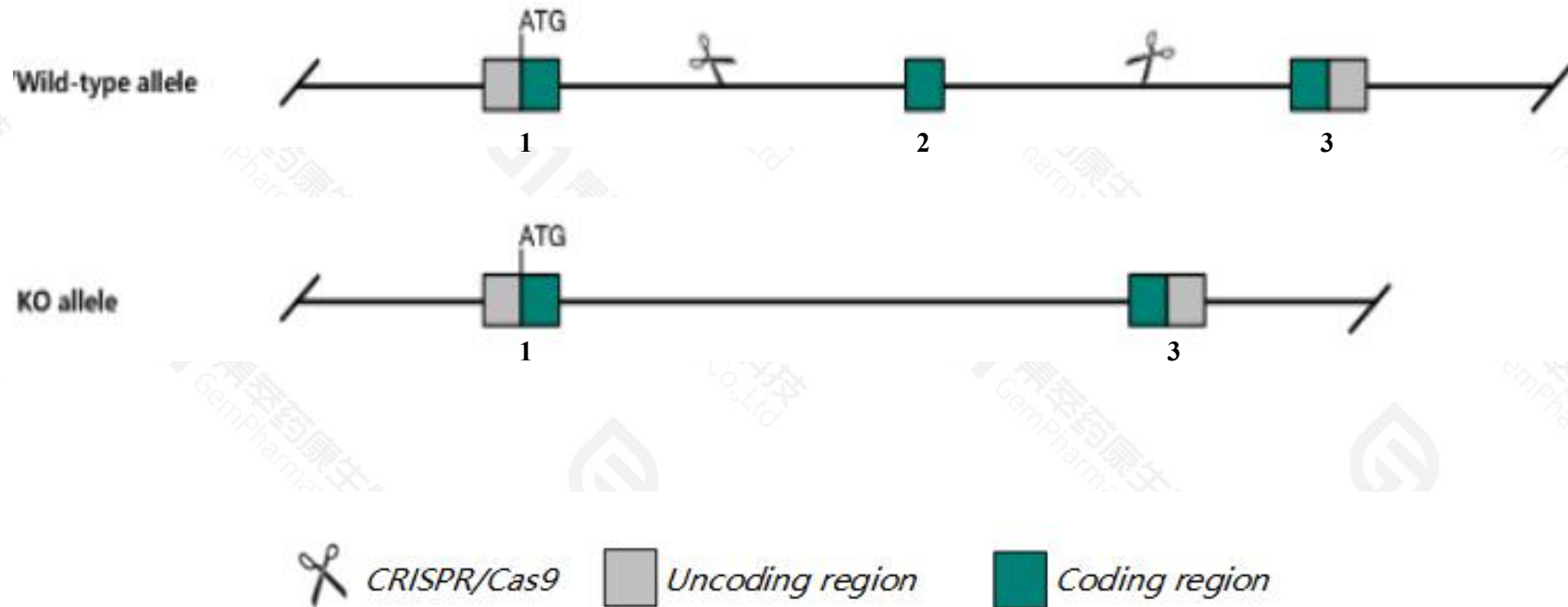
**Strain background**

**BALB/cJGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Il17a* gene. The schematic diagram is as follows:



- The *Il17a* gene has 1 transcript. According to the structure of *Il17a* gene, exon2 of *Il17a*-MGP\_BALBcJ\_T0018992.1 transcript is recommended as the knockout region. The region contains 212bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Il17a* gene. The brief process is as follows: CRISPR/Cas9 system were microinjected into the fertilized eggs of BALB/cJGpt 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 BALB/cJGpt mice.



- According to the existing MGI data, homozygotes for a targeted null mutation exhibit reduced contact, delayed-type and airway hypersensitivity responses and impaired T-dependent antibody production.
- The *Il17a* gene is located on the Chr1. 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.

## Il17a interleukin 17A [Mus musculus (house mouse)]

Gene ID: 16171, updated on 7-Mar-2021

### Summary



**Official Symbol** Il17a provided by [MGI](#)

**Official Full Name** interleukin 17A provided by [MGI](#)

**Primary source** [MGI:MGI:107364](#)

**See related** [Ensembl:ENSMUSG00000025929](#)

**Gene type** protein coding

**RefSeq status** REVIEWED

**Organism** [Mus musculus](#)

**Lineage** Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

**Also known as** Ctl, Ctla, Ctla-8, Ctla8, IL-, IL-17, IL-17A, Il, Il17

**Summary** This gene encodes a pro-inflammatory cytokine that is a member of the interleukin-17 family. The encoded protein plays a central role in host defense against diverse pathogens. The encoded protein is produced by activated T-cells and certain cell types of innate immune system. The active protein functions as either a homodimer with other interleukin-17 family members and signals through the interleukin-17 receptor to induce inflammatory cytokine production. Aberrant expression of this gene is associated with autoinflammatory diseases including rheumatoid arthritis, psoriasis and multiple sclerosis. [provided by RefSeq, Sep 2015]

**Expression** Low expression observed in reference dataset [See more](#)

**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

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

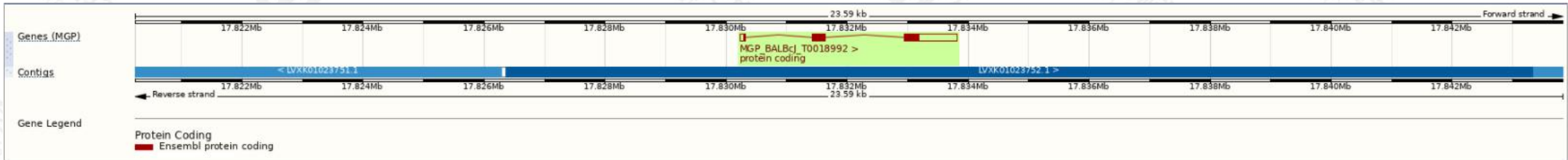
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt Match	Flags
-	<a href="#">MGP_BALBcJ_T0018992.1</a>	1171	<a href="#">158aa</a>	<span style="color: red;"> </span> Protein coding	<a href="#">CCDS14842</a>	<a href="#">Q544E6</a> <a href="#">Q62386</a>	-

The strategy is based on the design of *Il17a*-MGP\_BALBcJ\_T0018992.1 transcript, the transcription is shown below:



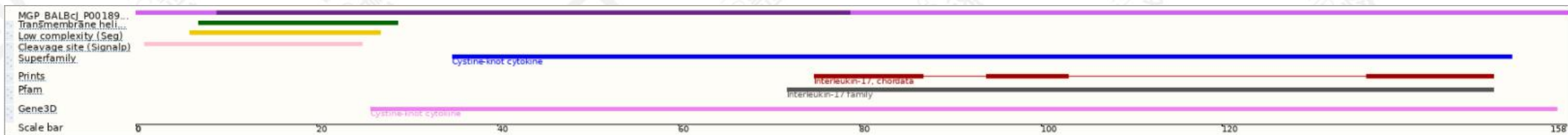


# Genomic location distribution

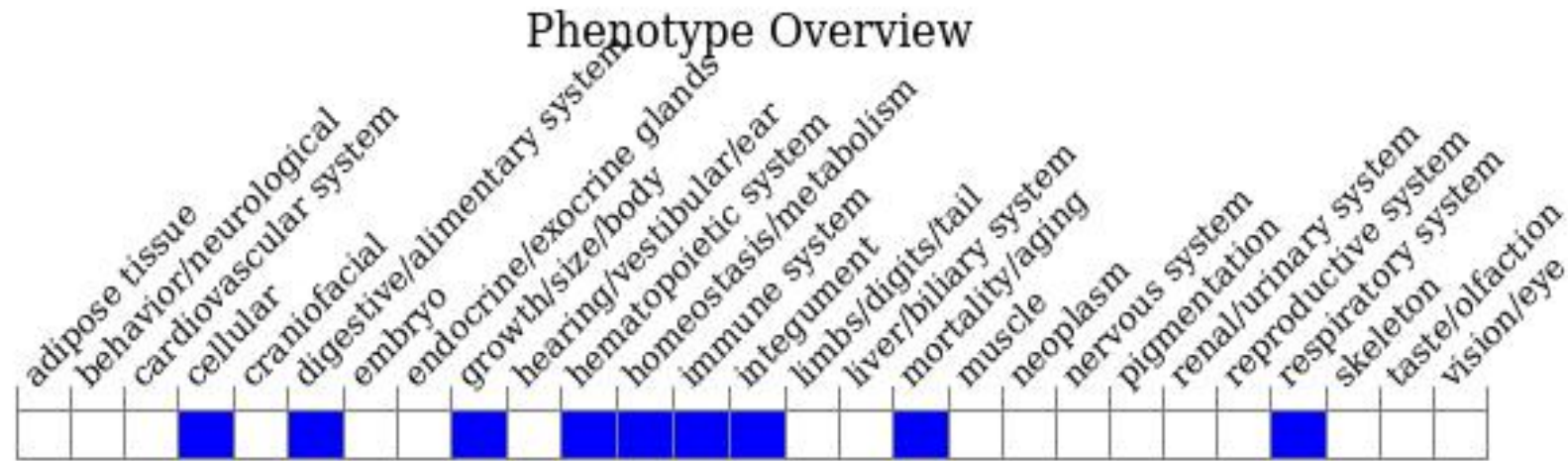




# Protein domain



# 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, homozygotes for a targeted null mutation exhibit reduced contact, delayed-type and airway hypersensitivity responses and impaired T-dependent antibody production.

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