

# *Cish* Cas9-KO Strategy

**Designer: Shilei Zhu**

**Reviewer: Lingyan Wu**

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# Project Overview

**Project Name**

*Cish*

**Project type**

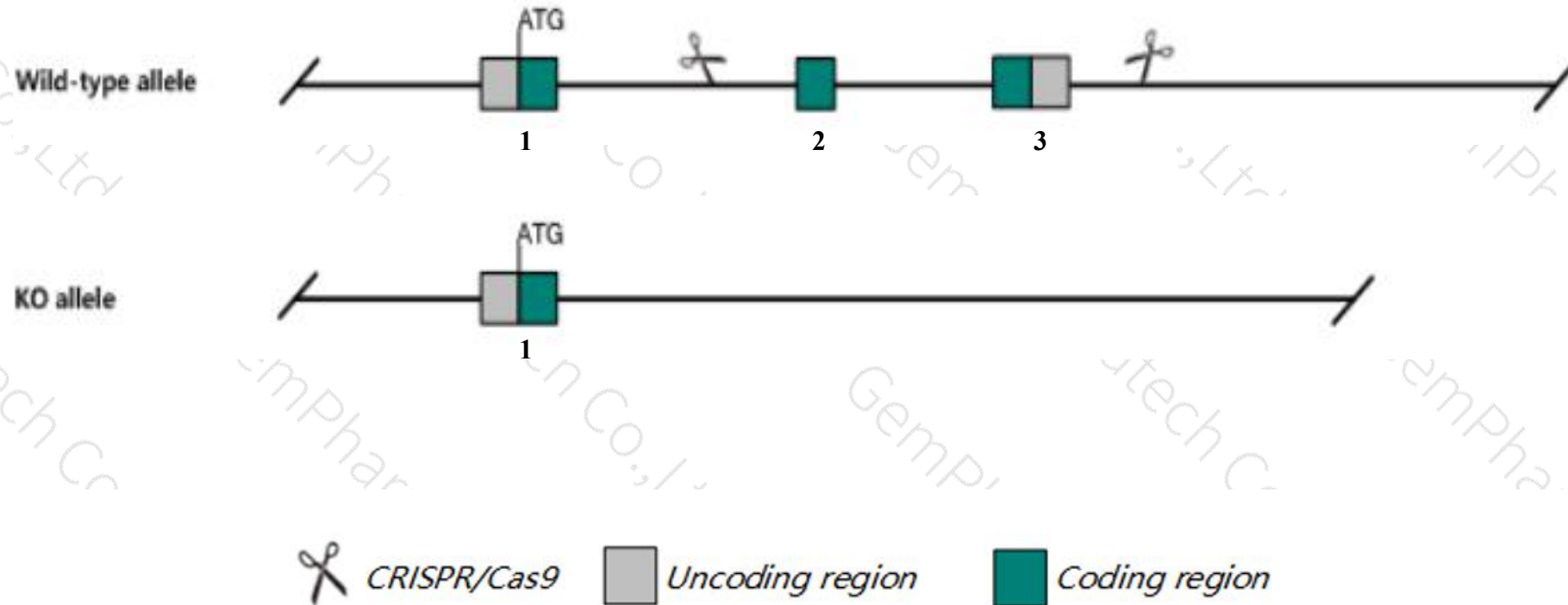
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

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



- The *Cish* gene has 5 transcripts. According to the structure of *Cish* gene, exon2-exon3 of *Cish*-201(ENSMUST00000085102.5) transcript is recommended as the knockout region. The region contains 754bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cish* gene. The brief process is as follows: CRISPR/Cas9 system 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.

- According to the existing MGI data, mice homozygous for a knock-out allele exhibit enhanced Th2 and Th9 differentiation and allergic airway inflammation.
- The *Cish* gene is located on the Chr9. 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.
- The deleted area contains GM17040-202, 5 'loxP will be placed on E1 of Cish-202(processed transcript).



# Gene information (NCBI)

## Cish cytokine inducible SH2-containing protein [Mus musculus (house mouse)]

Gene ID: 12700, updated on 13-Mar-2020

### Summary



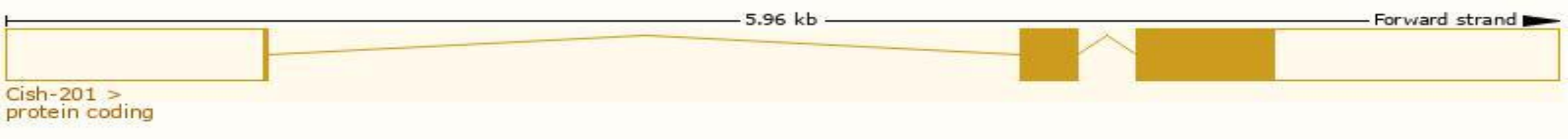
<b>Official Symbol</b>	Cish provided by <a href="#">MGI</a>
<b>Official Full Name</b>	cytokine inducible SH2-containing protein provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:103159</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000032578</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	CIS-1, CIS1, Cis, F17, F23, SOCS
<b>Expression</b>	Broad expression in heart adult (RPKM 25.8), lung adult (RPKM 20.1) and 24 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

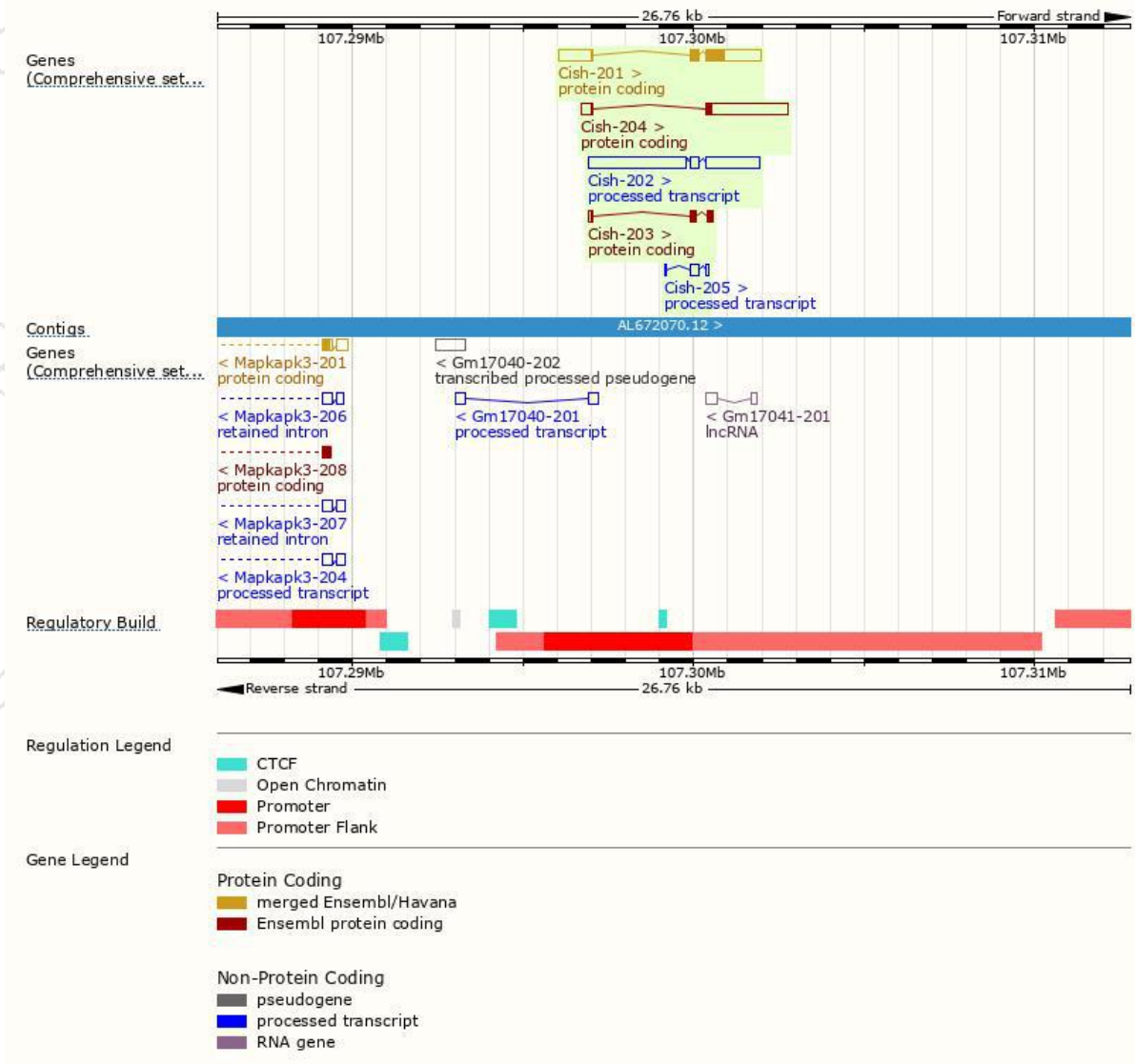
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cish-201	<a href="#">ENSMUST00000085102.5</a>	2850	<a href="#">257aa</a>	Protein coding	<a href="#">CCDS40761</a>	<a href="#">Q62225</a>	TSL:1 GENCODE basic APPRIS P1
Cish-204	<a href="#">ENSMUST00000168260.1</a>	2742	<a href="#">62aa</a>	Protein coding	-	<a href="#">E9Q273</a>	TSL:1 GENCODE basic
Cish-203	<a href="#">ENSMUST00000167072.1</a>	411	<a href="#">102aa</a>	Protein coding	-	<a href="#">E9Q2F4</a>	CDS 3' incomplete TSL:5
Cish-202	<a href="#">ENSMUST00000165664.1</a>	4697	No protein	Processed transcript	-	-	TSL:5
Cish-205	<a href="#">ENSMUST00000171568.1</a>	371	No protein	Processed transcript	-	-	TSL:3

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

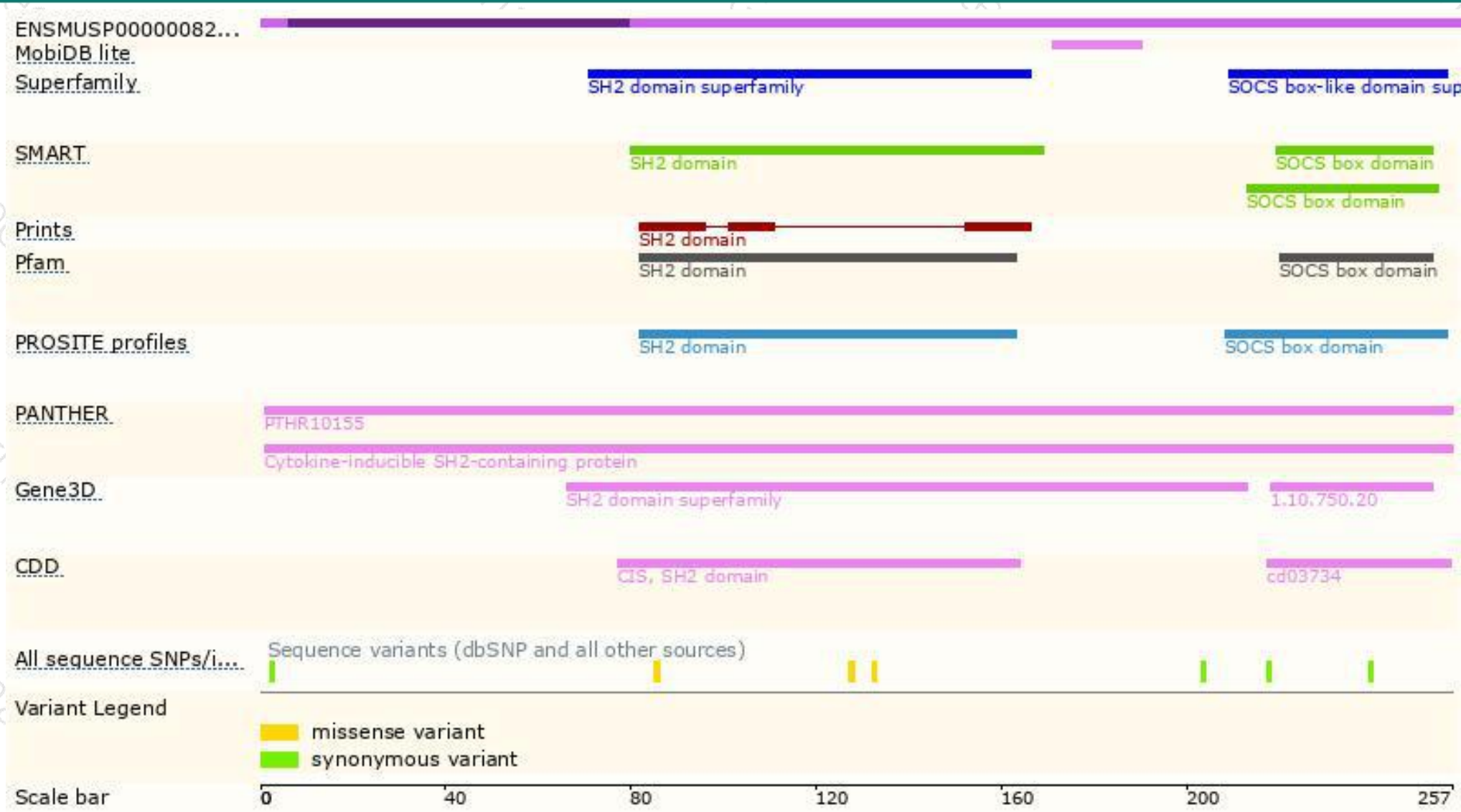


# Genomic location distribution

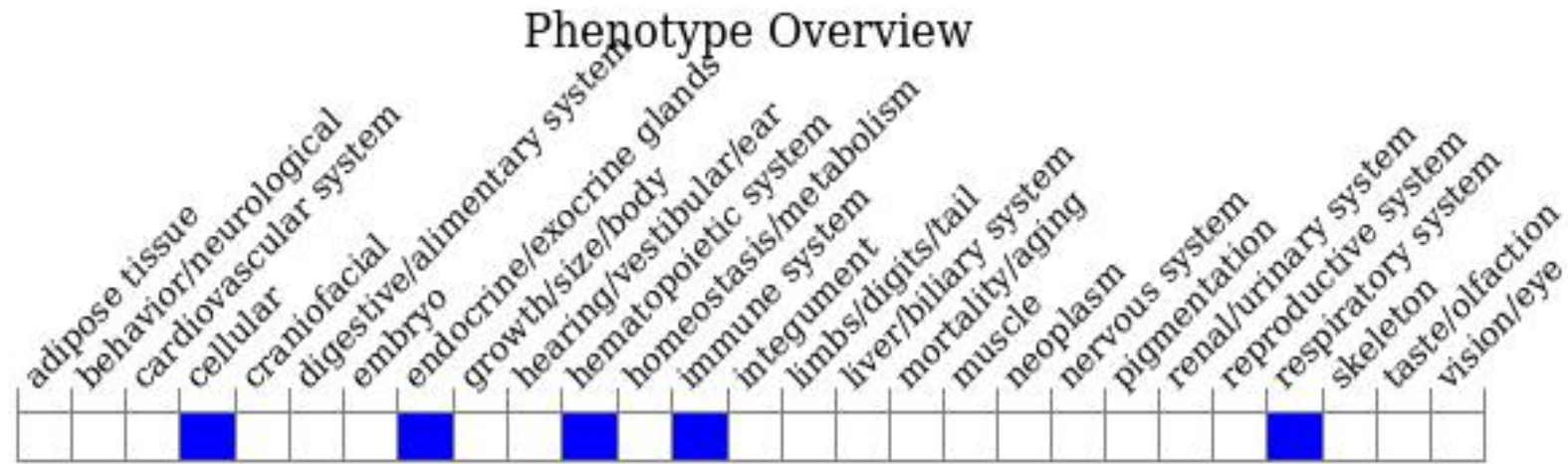




# Protein domain



# 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, mice homozygous for a knock-out allele exhibit enhanced Th2 and Th9 differentiation and allergic airway inflammation.

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

Tel: 400-9660890

