

# ***Il1b*** Cas9-KO Strategy

Designer: Xueting Zhang

# Project Overview

**Project Name**

***Il1b***

**Project type**

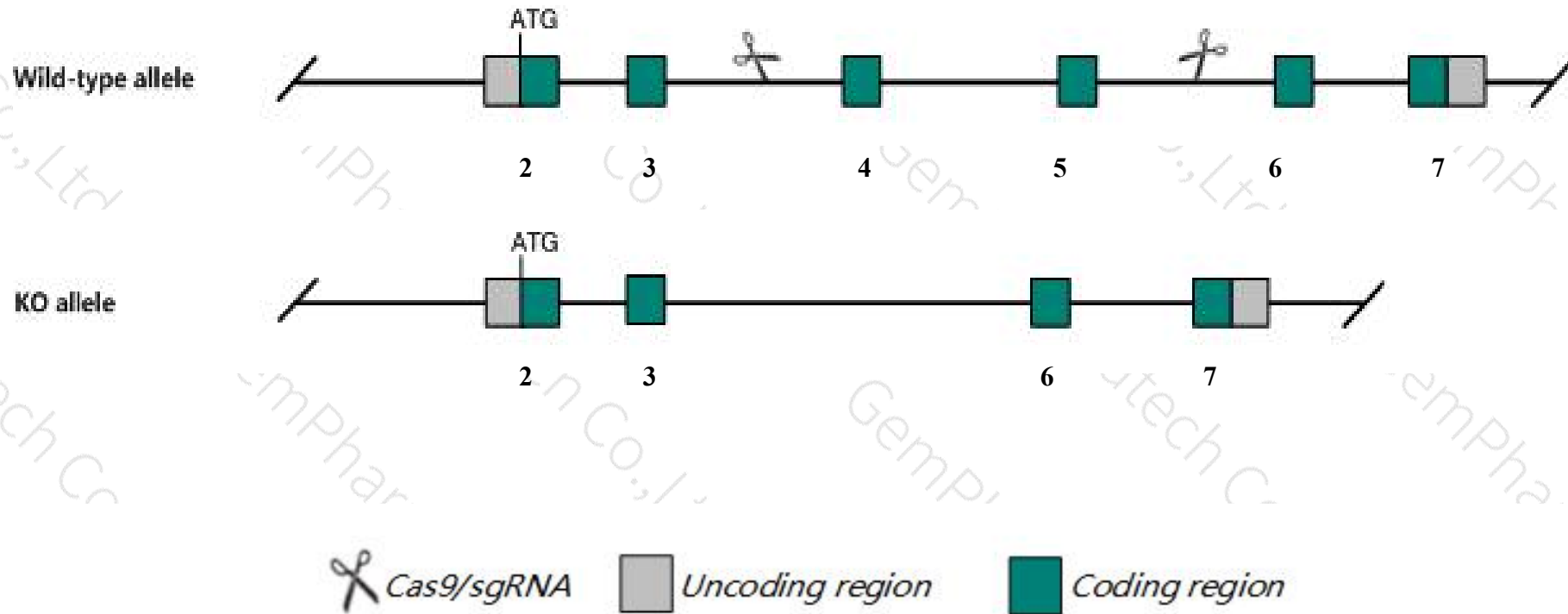
**Cas9-KO**

**Strain background**

**C57BL/6J**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Il1b* gene. The schematic diagram is as follows:



- The *Il1b* gene has 3 transcripts. According to the structure of *Il1b* gene, exon4-exon5 of *Il1b*-201 (ENSMUST00000028881.13) transcript is recommended as the knockout region. The region contains 373bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Il1b* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

- According to the existing MGI data, Homozygous null mutants show impaired contact hypersensitivity and reduced acute-phase inflammatory response. Lung tumors and metastases of B16 melanoma do not occur in null mutant mice, suggesting inability to support tumor invasiveness and angiogenesis.
- The *Il1b* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.



# Gene information (NCBI)

## Il1b interleukin 1 beta [Mus musculus (house mouse)]

Gene ID: 16176, updated on 9-Apr-2019

### Summary



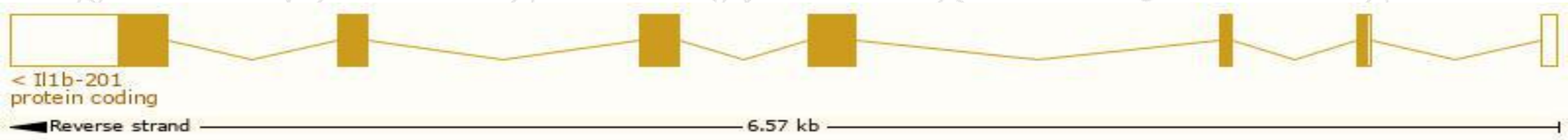
<b>Official Symbol</b>	Il1b provided by <a href="#">MGI</a>
<b>Official Full Name</b>	interleukin 1 beta provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:96543</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG000000027398</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	REVIEWED
<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>	IL-1beta, Il-1b
<b>Summary</b>	The protein encoded by this gene is a member of the interleukin 1 cytokine family. This cytokine is produced by activated macrophages as a proprotein, which is proteolytically processed to its active form by caspase 1. The encoded protein plays a role in thymocyte proliferation and is involved in the inflammatory response. [provided by RefSeq, Aug 2015]
<b>Expression</b>	Broad expression in spleen adult (RPKM 1.3), lung adult (RPKM 1.0) and 19 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

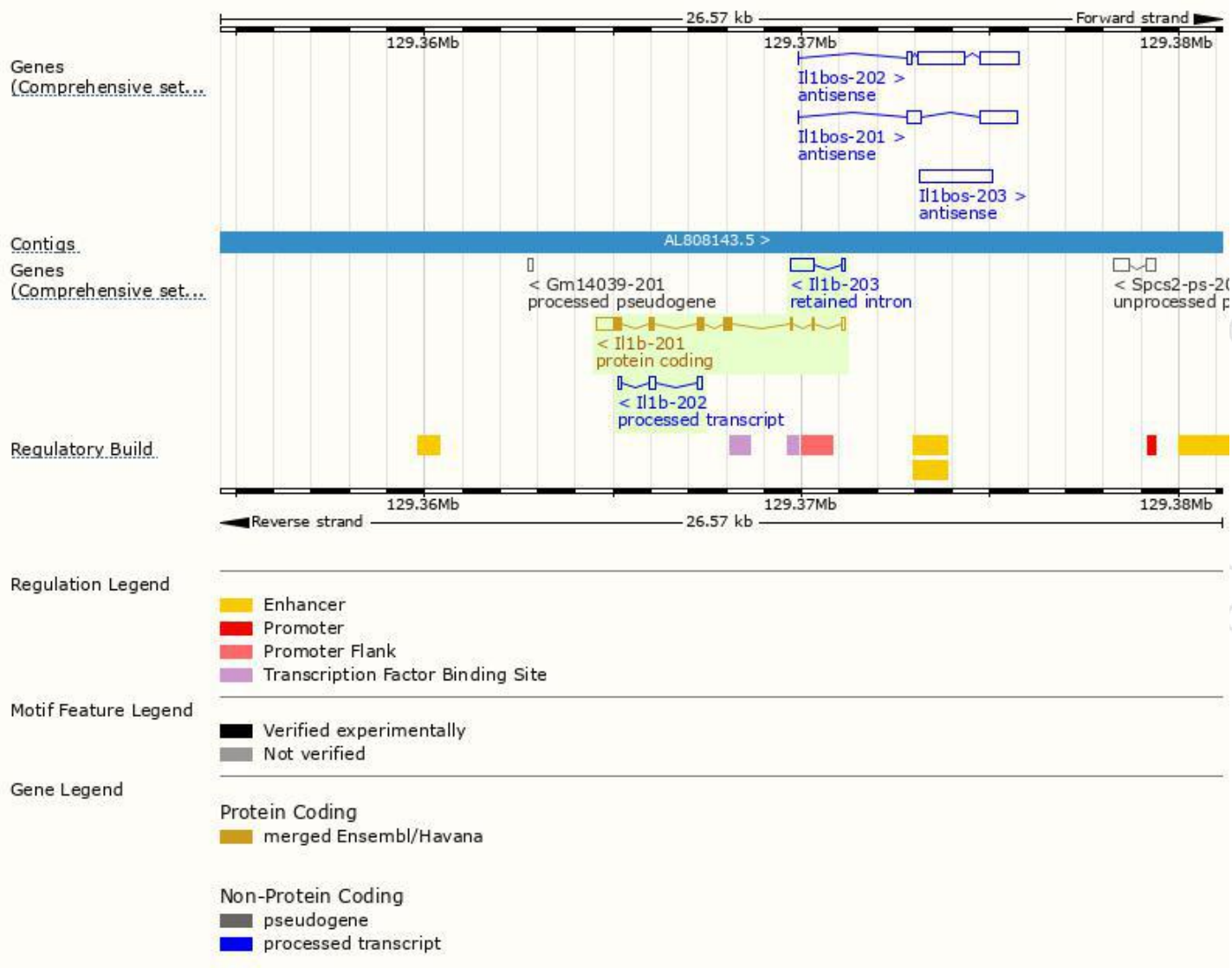
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Il1b-201	<a href="#">ENSMUST00000028881.13</a>	1356	<a href="#">269aa</a>	Protein coding	<a href="#">CCDS16726</a>	<a href="#">P10749</a>	TSL:1 GENCODE basic APPRIS P1
Il1b-202	<a href="#">ENSMUST00000141979.1</a>	379	No protein	Processed transcript	-	-	TSL:1
Il1b-203	<a href="#">ENSMUST00000155994.1</a>	689	No protein	Retained intron	-	-	TSL:2

The strategy is based on the design of *Il1b-201* transcript,The transcription is shown below

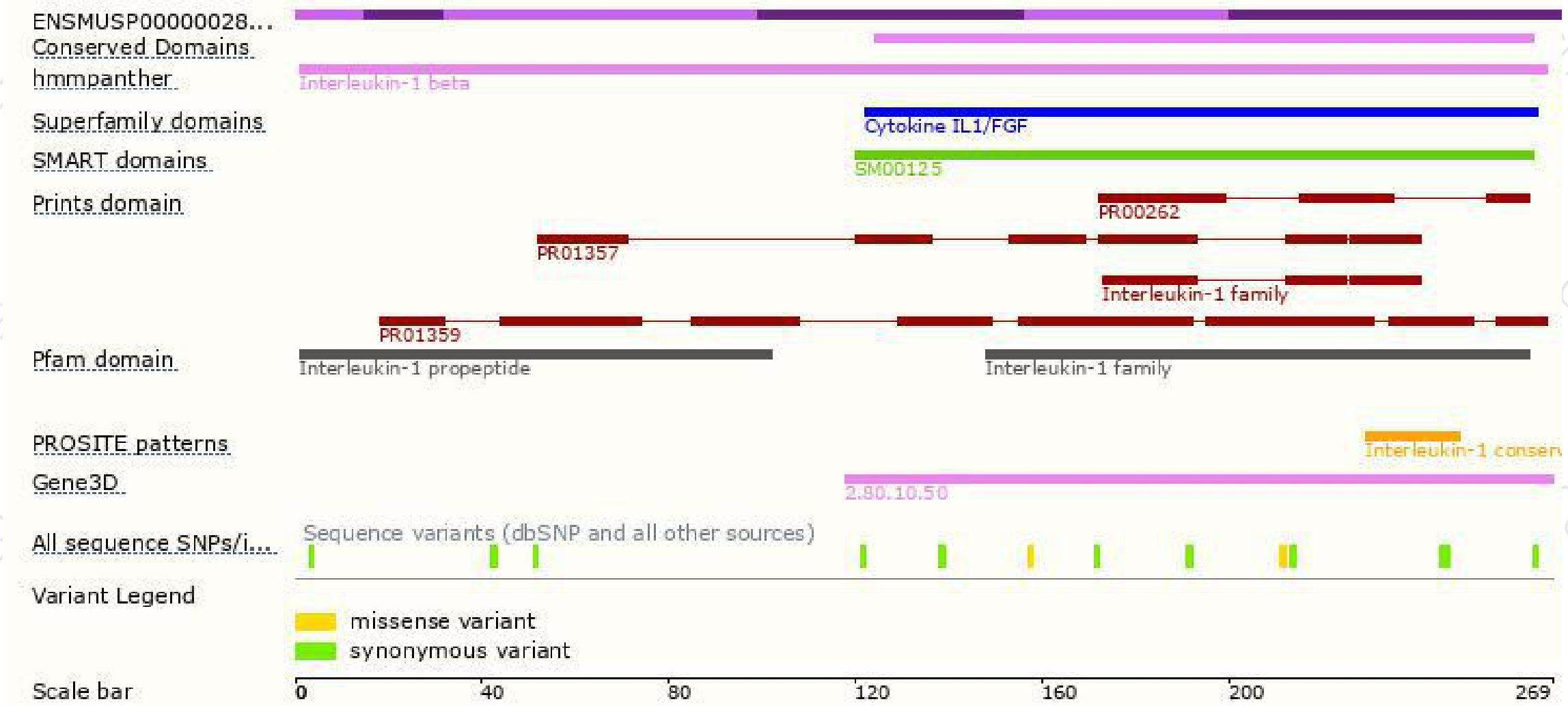


# Genomic location distribution

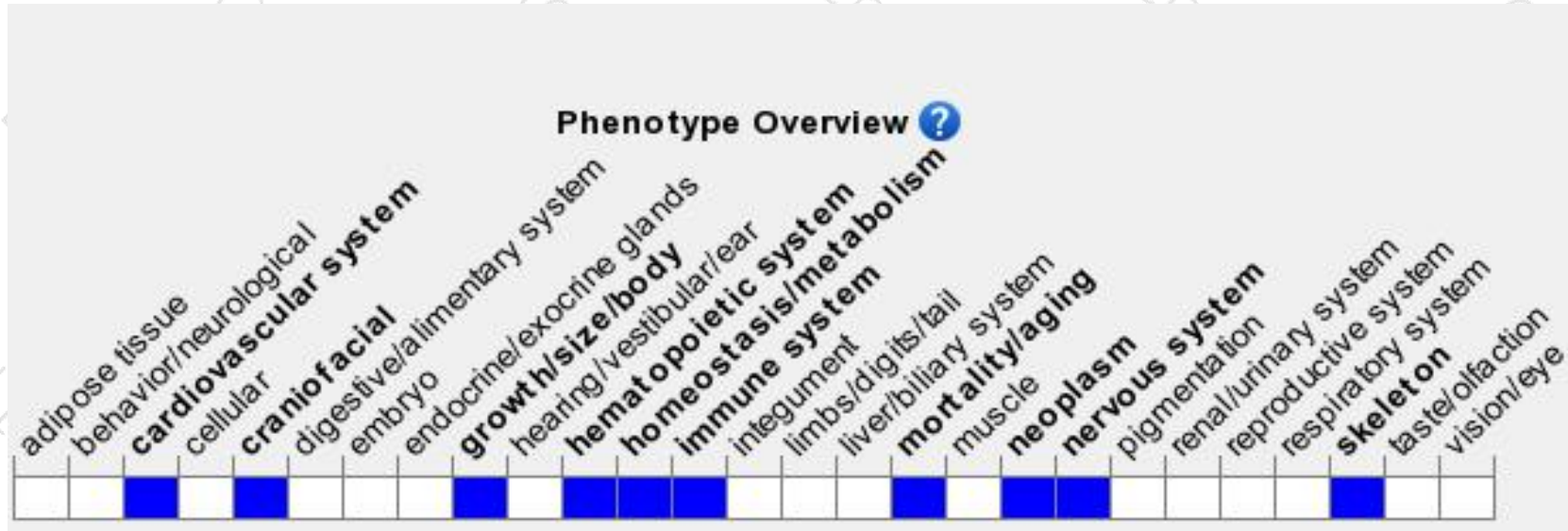




# Protein domain



# 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 show impaired contact hypersensitivity and reduced acute-phase inflammatory response. Lung tumors and metastases of B16 melanoma do not occur in null mutant mice, suggesting inability to support tumor invasiveness and angiogenesis.

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

Tel: 025-5864 1534

