

# ***Il1a*** Cas9-KO Strategy

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# Project Overview

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**Project Name**

***Il1a***

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**Project type**

**Cas9-KO**

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**Strain background**

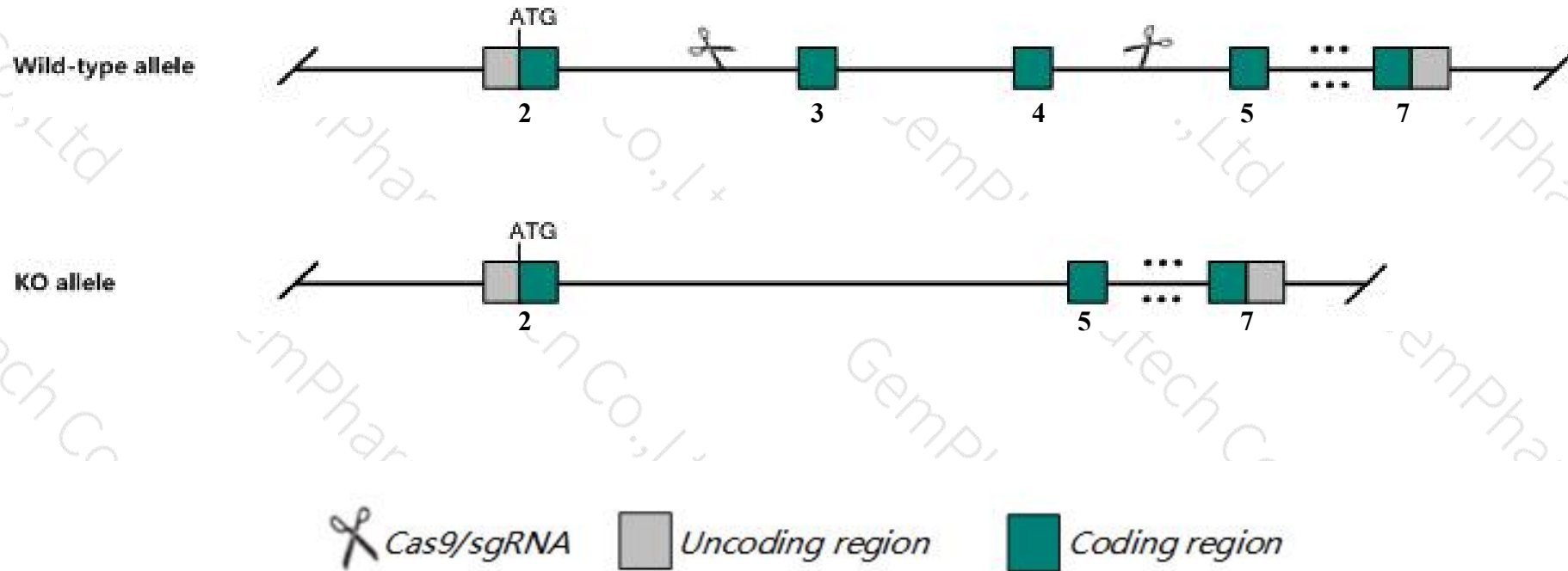
**C57BL/6J**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Il1a* gene. The schematic diagram is as follows:





- The *Il1a* gene has 1 transcript. According to the structure of *Il1a* gene, exon3-exon4 of *Il1a-201* (ENSMUST00000028882.1) transcript is recommended as the knockout region. The region contains 278bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Il1a* 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, Mice homozygous for a knock-out allele exhibit abnormal tumor vascularization, decreased metastatic potential, and decreased interleukin-1 beta secretion.
- The *Il1a* gene coincides with the Gm14023 gene, and the exon4 and exon5 of the Gm14023 gene are deleted at the same time, and it is uncertain whether the phenotype of the mouse is caused by knockout of which gene.
- The *Il1a* 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)

## Il1a interleukin 1 alpha [Mus musculus (house mouse)]

Gene ID: 16175, updated on 9-Apr-2019

### Summary



<b>Official Symbol</b>	Il1a provided by <a href="#">MGI</a>
<b>Official Full Name</b>	interleukin 1 alpha provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:96542</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG000000027399</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>	Il-1a
<b>Expression</b>	Broad expression in liver E18 (RPKM 1.2), liver E14 (RPKM 1.1) and 20 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

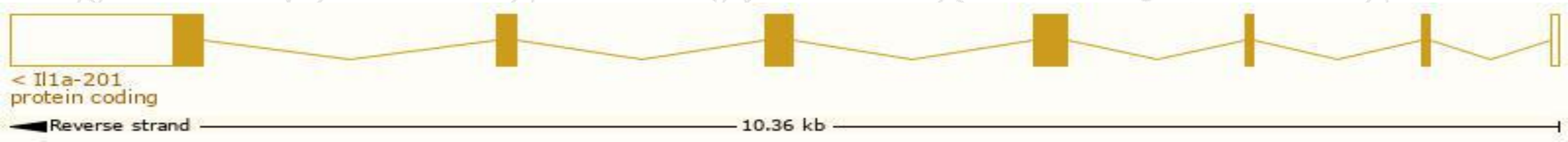


# Transcript information (Ensembl)

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

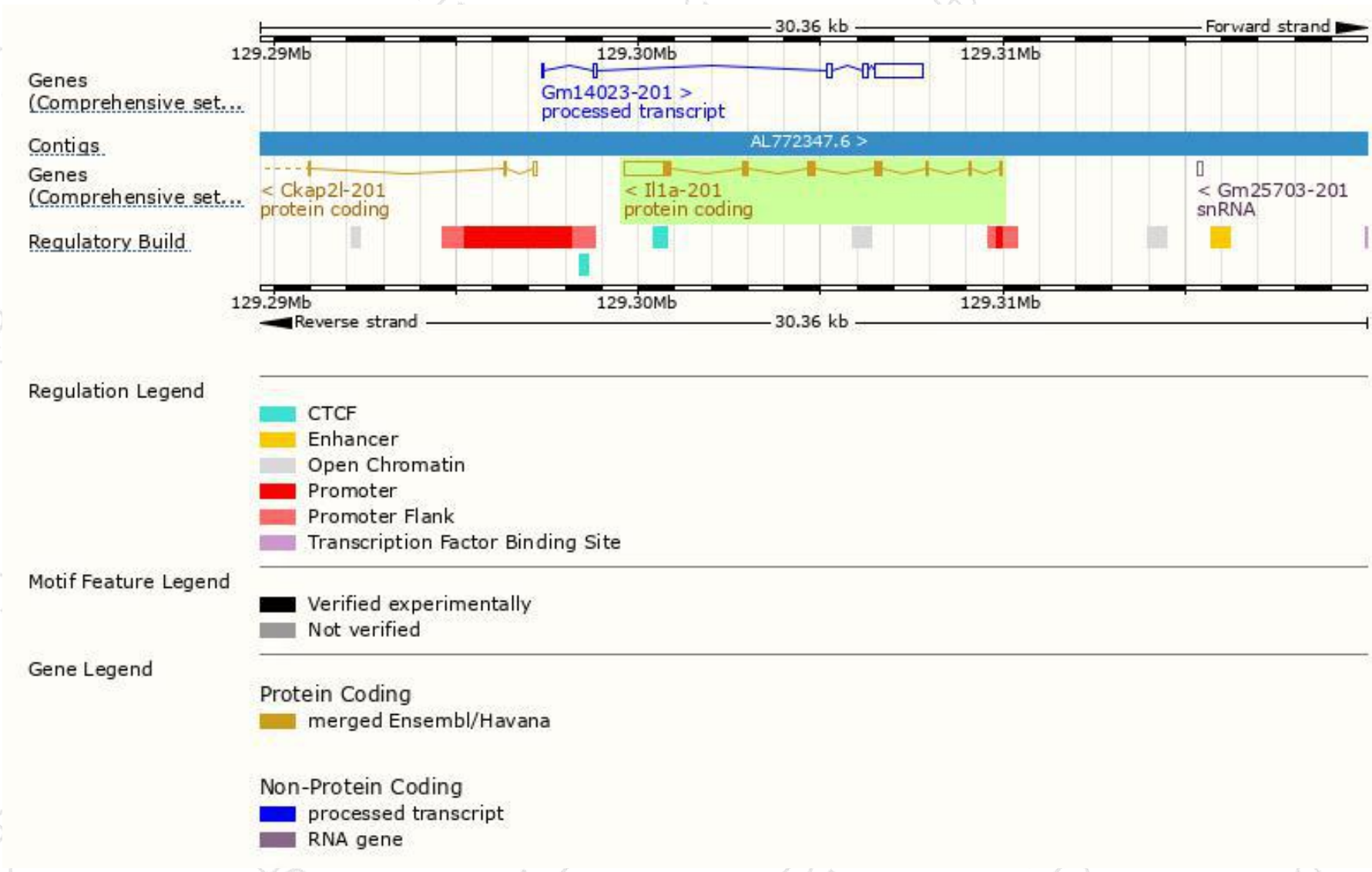
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Il1a-201	<a href="#">ENSMUST00000028882.1</a>	1974	<a href="#">270aa</a>	Protein coding	<a href="#">CCDS16725</a>	<a href="#">P01582</a> <a href="#">Q3U0Y6</a>	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Il1a-201* transcript, The transcription is shown below



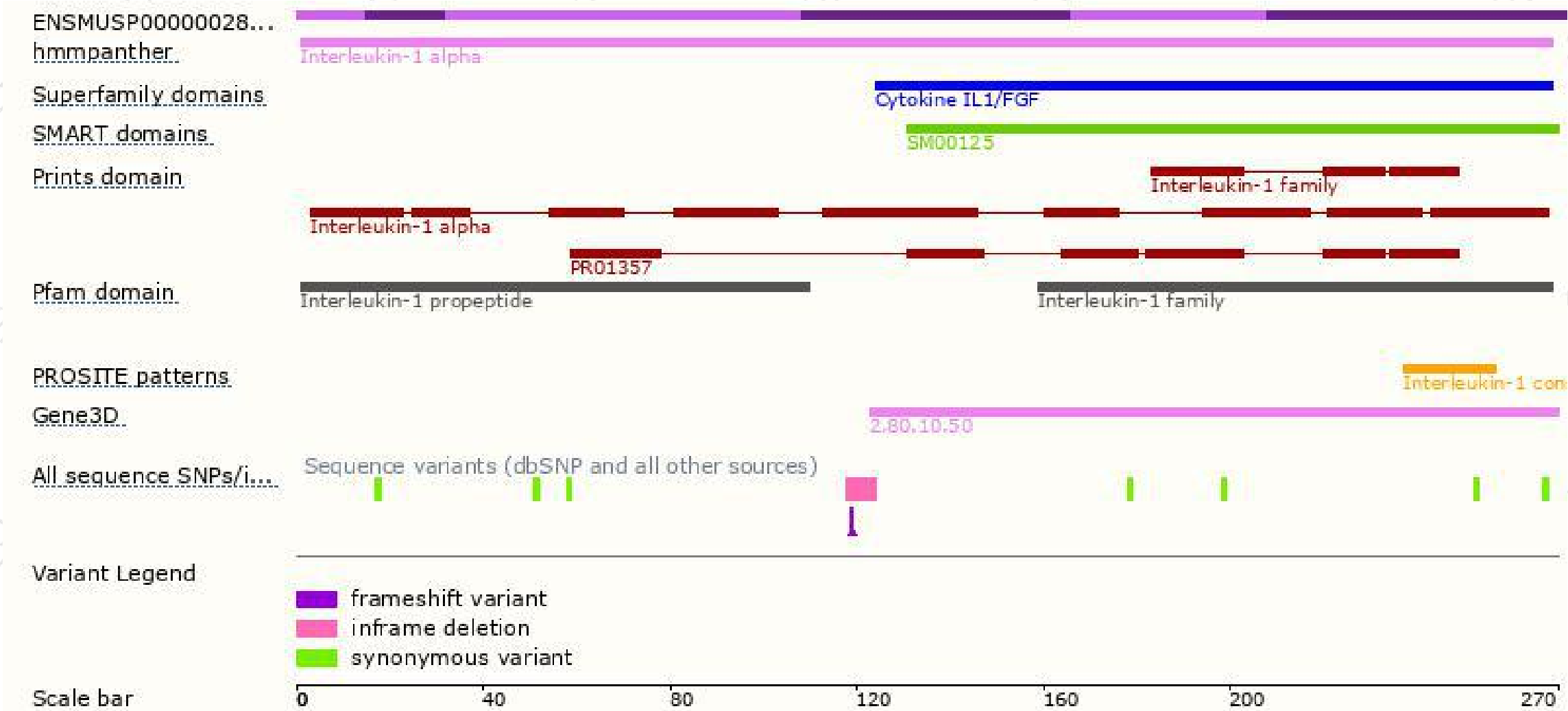


# Genomic location distribution



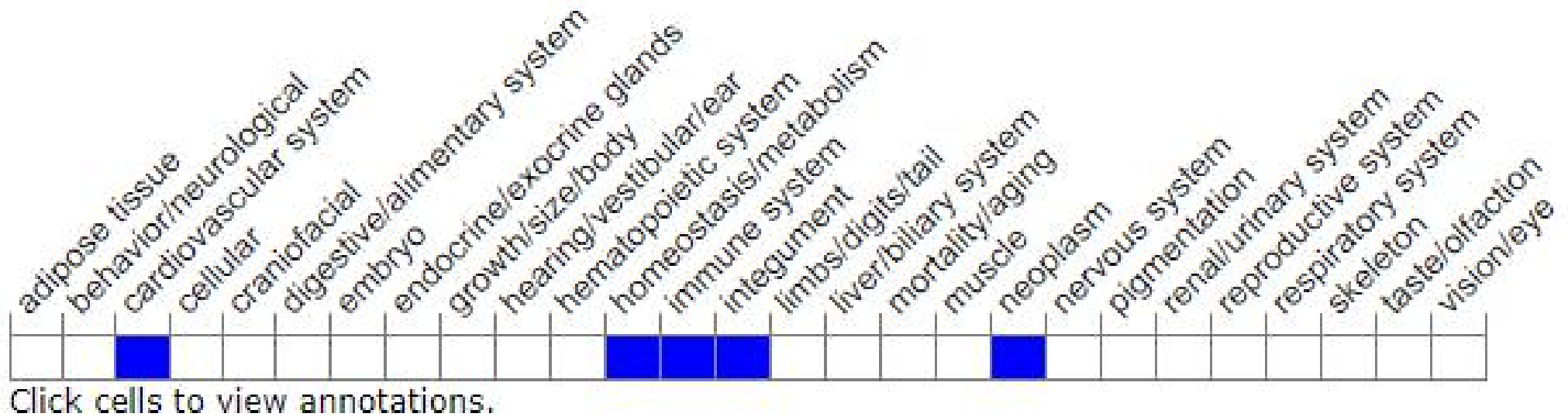


# Protein domain





# 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 abnormal tumor vascularization, decreased metastatic potential, and decreased interleukin-1 beta secretion.



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

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