

# ***Tec Cas9-CKO Strategy***

**Designer:**

**Yang Zeng**

**Reviewer:**

**Huimin su**

**Design Date:**

**2019-11-21**

# Project Overview

**Project Name**

***Tec***

**Project type**

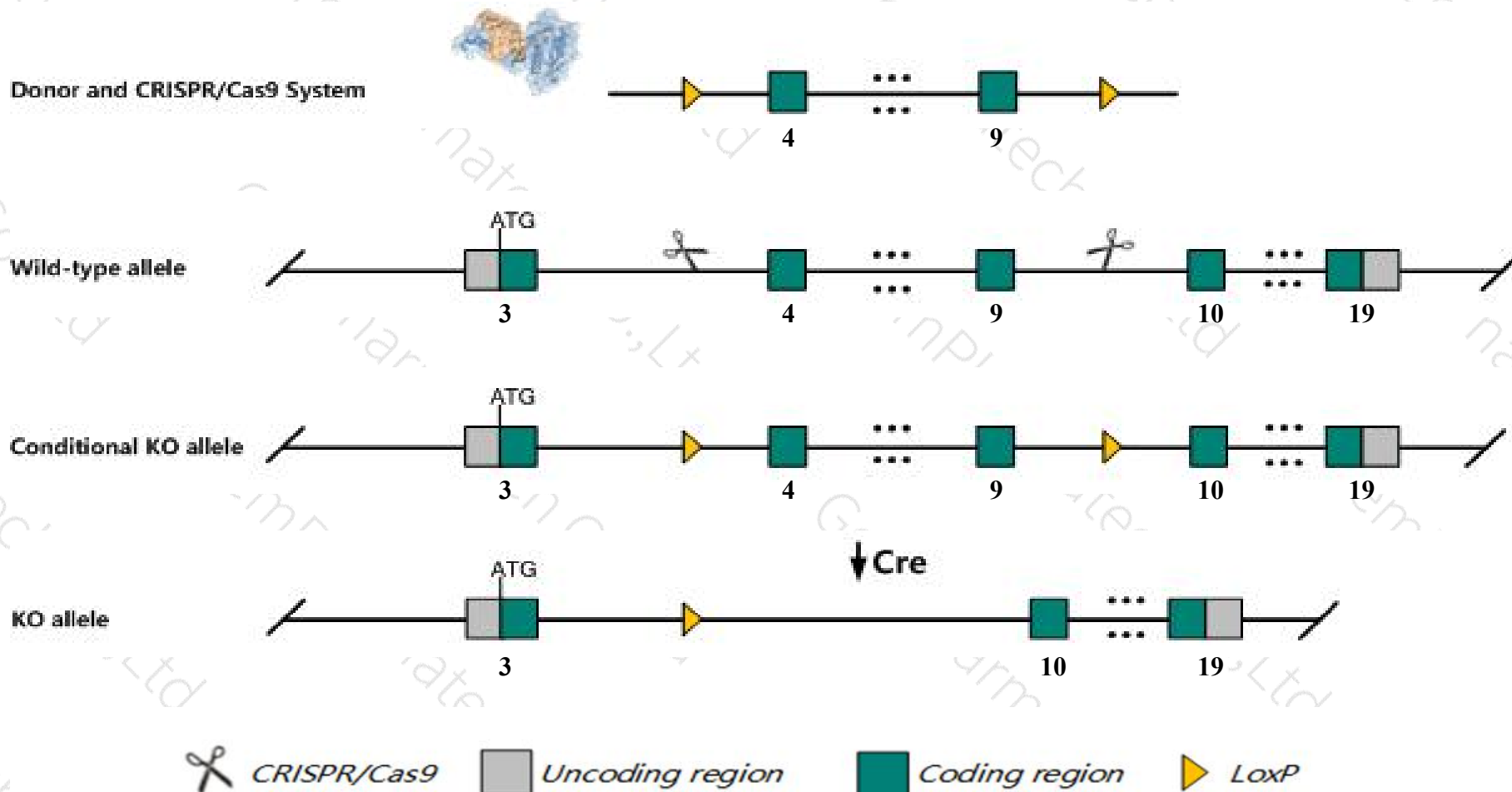
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

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



- The *Tec* gene has 9 transcripts. According to the structure of *Tec* gene, exon4-exon9 of *Tec-201* (ENSMUST00000071944.12) transcript is recommended as the knockout region. The region contains 596bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Tec* 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.

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit a minor reduction in platelet aggregation in response to threshold concentrations of collagen-related peptide or collagen.
- Transcript *Tec*-209 CDS 3' is incomplete affected, whether it will be affected is unknown.
- The *Tec* gene is located on the Chr5. 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.



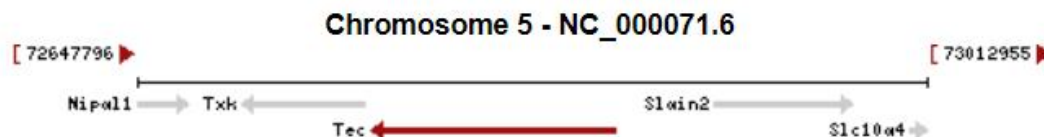
# Gene information (NCBI)

## Tec tec protein tyrosine kinase [ *Mus musculus* (house mouse) ]

Gene ID: 21682, updated on 12-Aug-2019

### Summary

<b>Official Symbol</b>	Tec provided by <a href="#">MGI</a>
<b>Official Full Name</b>	tec protein tyrosine kinase provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:98662</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000029217</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>Expression</b>	Ubiquitous expression in placenta adult (RPKM 4.6), spleen adult (RPKM 4.0) and 27 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

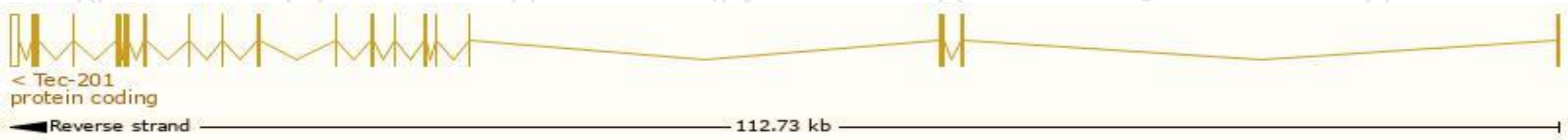


# Transcript information (Ensembl)

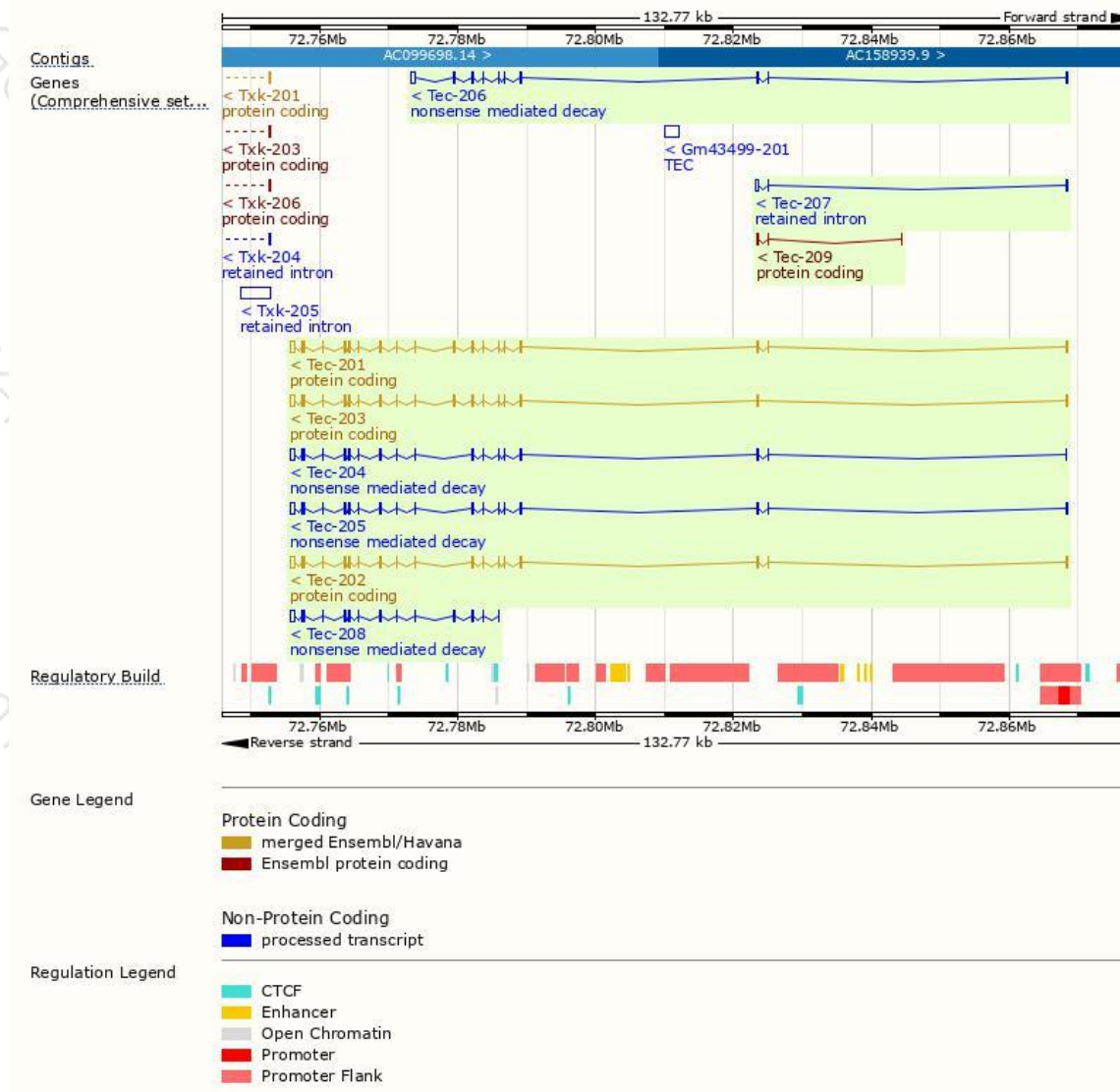
The gene has 9 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tec-201	<a href="#">ENSMUST00000071944.12</a>	2659	<a href="#">630aa</a>	Protein coding	<a href="#">CCDS51516</a>	<a href="#">Q3U436</a>	TSL:1 GENCODE basic APPRIS P1
Tec-203	<a href="#">ENSMUST00000113594.7</a>	2580	<a href="#">630aa</a>	Protein coding	<a href="#">CCDS51516</a>	<a href="#">Q3U436</a>	TSL:1 GENCODE basic APPRIS P1
Tec-202	<a href="#">ENSMUST00000073843.12</a>	2573	<a href="#">608aa</a>	Protein coding	<a href="#">CCDS51515</a>	<a href="#">Q8CFK4</a>	TSL:1 GENCODE basic
Tec-209	<a href="#">ENSMUST00000202547.1</a>	377	<a href="#">44aa</a>	Protein coding	-	<a href="#">A0A0J9YV29</a>	CDS 3' incomplete TSL:3
Tec-205	<a href="#">ENSMUST00000138842.7</a>	2569	<a href="#">100aa</a>	Nonsense mediated decay	-	<a href="#">A0A0R4J1V5</a>	TSL:1
Tec-204	<a href="#">ENSMUST00000126481.7</a>	2543	<a href="#">184aa</a>	Nonsense mediated decay	-	<a href="#">D6RJM5</a>	TSL:5
Tec-208	<a href="#">ENSMUST00000155342.7</a>	2117	<a href="#">68aa</a>	Nonsense mediated decay	-	<a href="#">F6W1T9</a>	CDS 5' incomplete TSL:2
Tec-206	<a href="#">ENSMUST00000149533.7</a>	1561	<a href="#">100aa</a>	Nonsense mediated decay	-	<a href="#">A0A0R4J1V5</a>	TSL:1
Tec-207	<a href="#">ENSMUST00000150193.1</a>	621	No protein	Retained intron	-	-	TSL:2

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

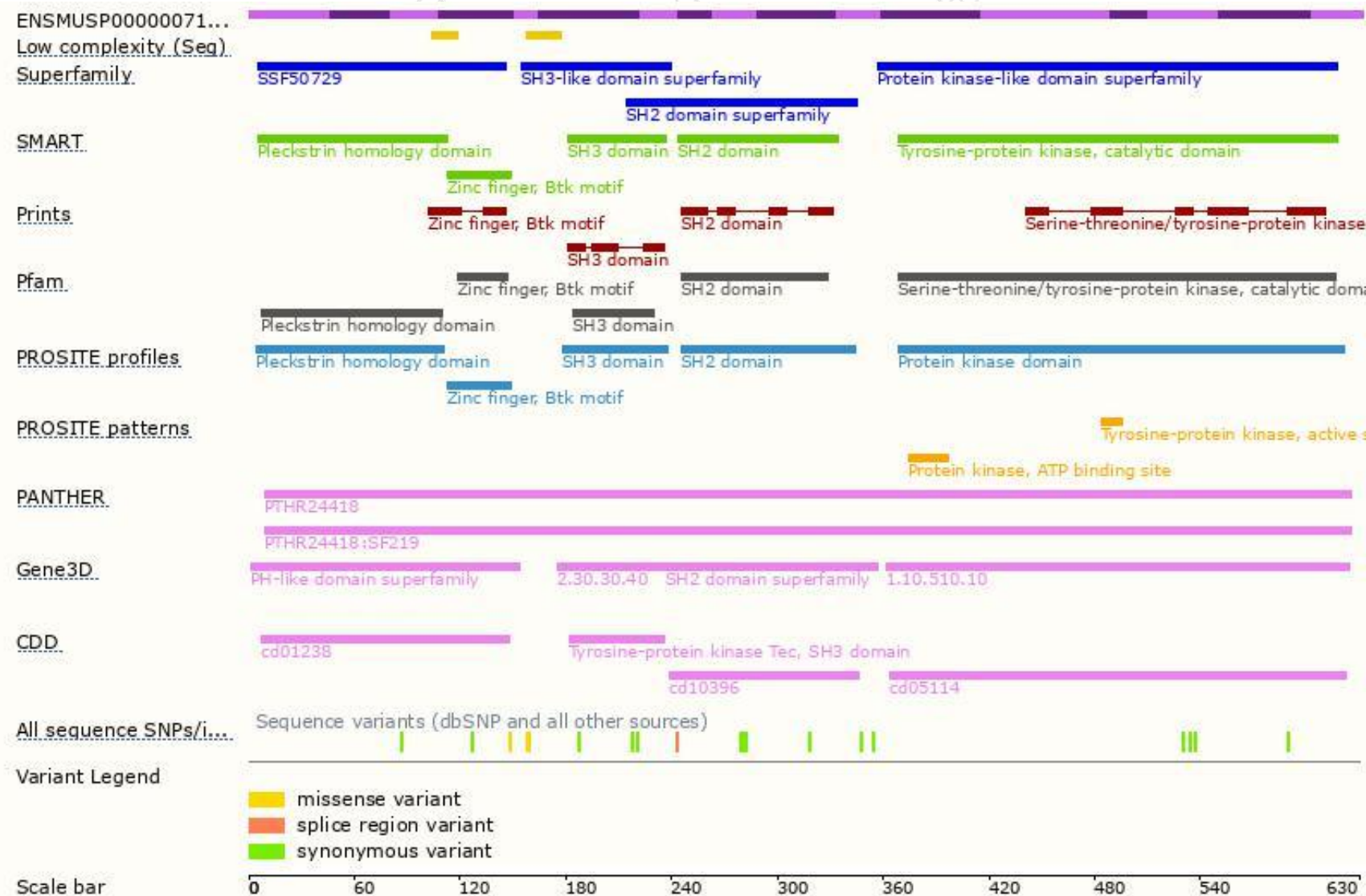


# Genomic location distribution

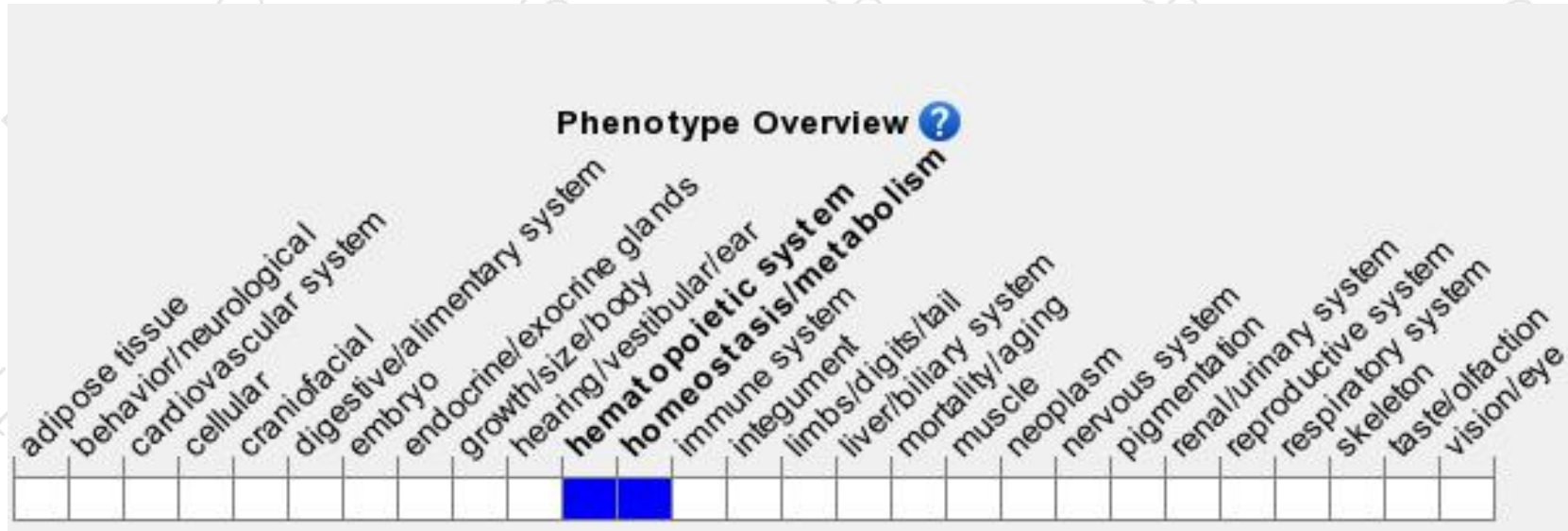




# Protein domain



# 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 a minor reduction in platelet aggregation in response to threshold concentrations of collagen-related peptide or collagen.

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

Tel: 400-9660890

