

Col4a1 Cas9-CKO Strategy

Designer: Shilei Zhu

Project Overview

Project Name

Col4a1

Project type

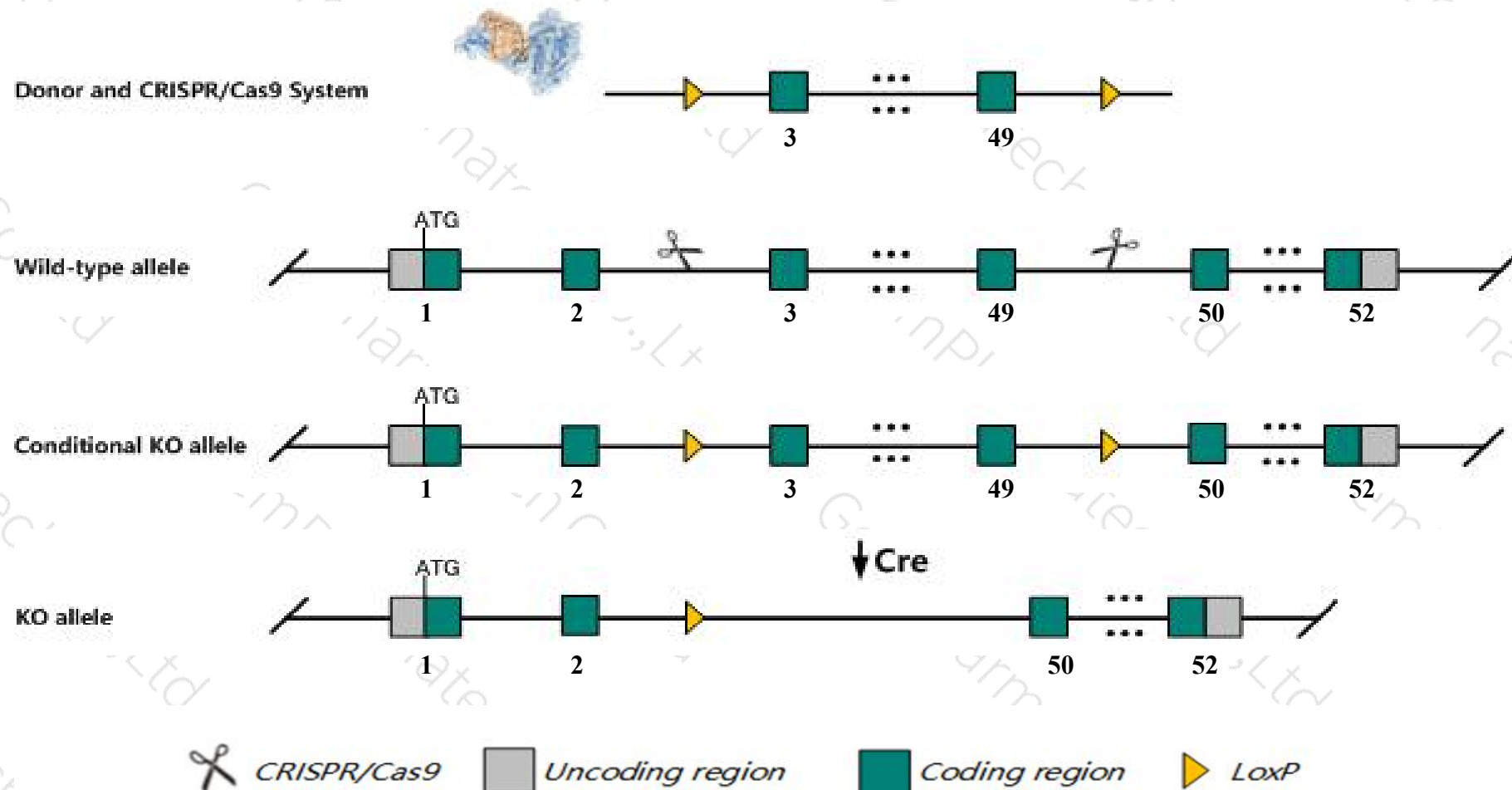
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Col4a1* gene. The schematic diagram is as follows:



- The *Col4a1* gene has 8 transcripts. According to the structure of *Col4a1* gene, exon3-exon49 of *Col4a1-201* (ENSMUST00000033898.9) transcript is recommended as the knockout region. The region contains 4496bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Col4a1* 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 with ENU induced alleles have various eye and vision defects and may show bruising at birth. Mice carrying the G498V mutation have renal glomerular defects that resolve within the first weeks of life, but show retinal tortuosity, muscular dystrophy, brain hemorrhages, and renal cysts as adults.
- The *Col4a1* gene is located on the Chr8. 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)

Col4a1 collagen, type IV, alpha 1 [Mus musculus (house mouse)]

Gene ID: 12826, updated on 27-Mar-2019

Summary



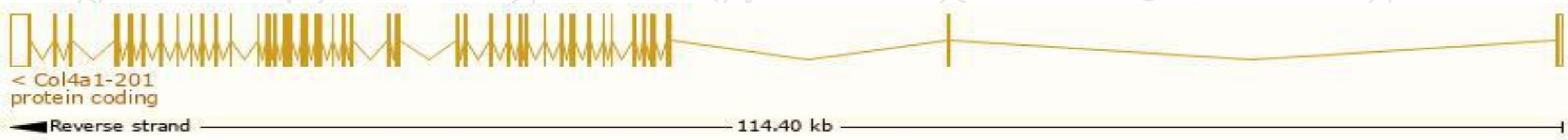
| | |
|---------------------------|--|
| Official Symbol | Col4a1 provided by MGI |
| Official Full Name | collagen, type IV, alpha 1 provided by MGI |
| Primary source | MGI:MGI:88454 |
| See related | Ensembl:ENSMUSG000000031502 |
| Gene type | protein coding |
| RefSeq status | REVIEWED |
| Organism | Mus musculus |
| Lineage | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| Also known as | Bru, Col4a-1, Raw, Svc |
| Summary | This gene encodes the alpha-1 subunit of the type IV collagens, an essential component of basement membranes. The encoded protein forms a triple helical heterotrimer comprised of two alpha-1 and one alpha-2 subunits that assembles into a type IV collagen network. This gene is located adjacent to the gene encoding alpha-2 subunit. Mice lacking both the alpha-1 and alpha-2 subunits of collagen IV die in utero due to structural deficiencies in the basement membranes and certain mutations in this gene cause perinatal cerebral hemorrhage and porencephaly. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Nov 2015] |
| Expression | Broad expression in subcutaneous fat pad adult (RPKM 160.4), lung adult (RPKM 107.4) and 21 other tissues See more |
| Orthologs | human all |

Transcript information (Ensembl)

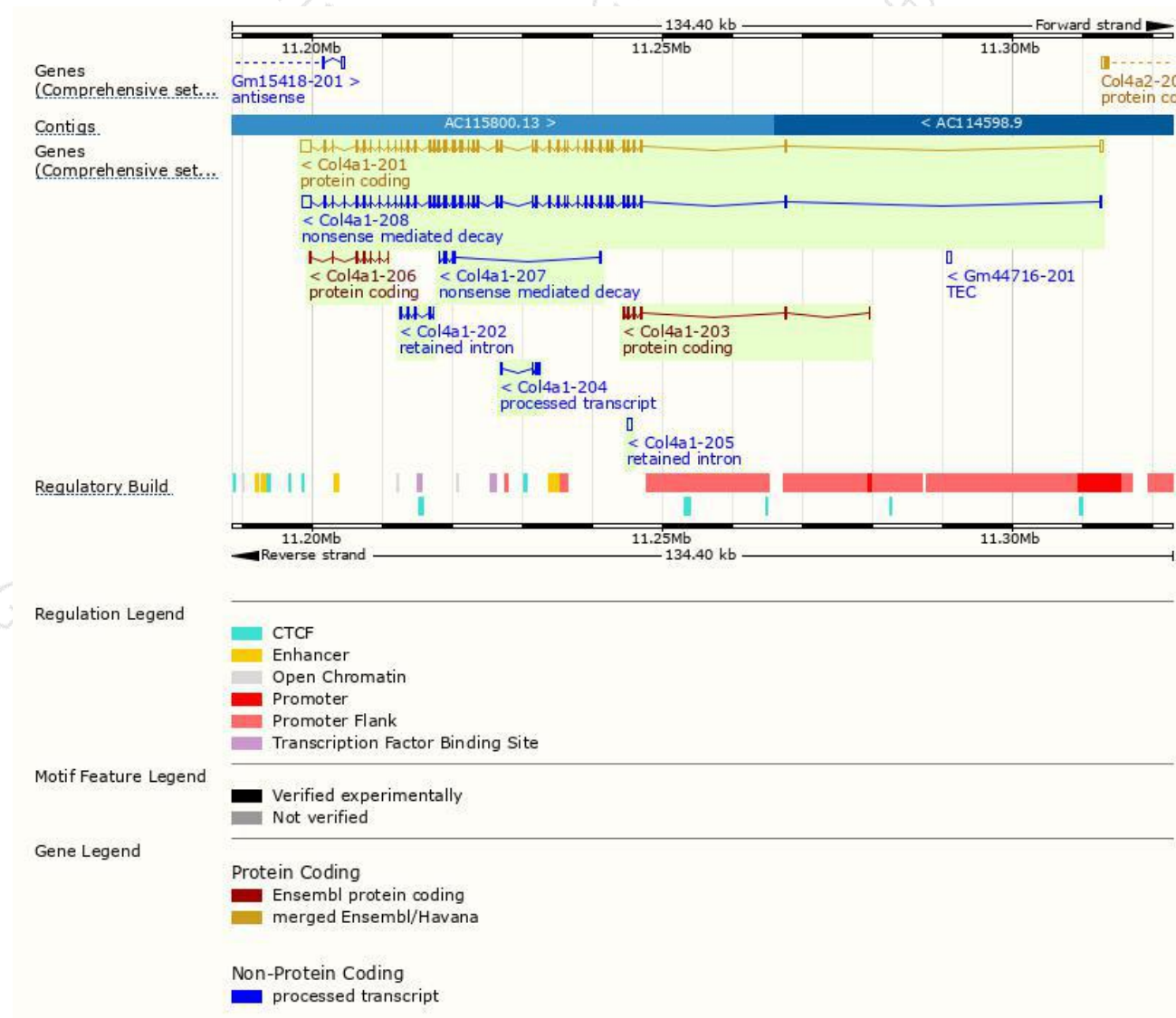
The gene has 8 transcripts,all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|------------|--------------------------------------|------|------------------------|-------------------------|---------------------------|----------------------------|-------------------------------|
| Col4a1-201 | ENSMUST00000033898.9 | 6615 | 1669aa | Protein coding | CCDS40219 | P02463 | TSL:1 GENCODE basic APPRIS P1 |
| Col4a1-206 | ENSMUST00000209598.1 | 1170 | 328aa | Protein coding | - | A0A1B0GRC0 | CDS 5' incomplete TSL:5 |
| Col4a1-203 | ENSMUST00000208095.1 | 545 | 133aa | Protein coding | - | A0A140LHU8 | CDS 3' incomplete TSL:3 |
| Col4a1-208 | ENSMUST00000209735.1 | 6487 | 1562aa | Nonsense mediated decay | - | A0A1B0GSI7 | TSL:2 |
| Col4a1-207 | ENSMUST00000209661.1 | 618 | 126aa | Nonsense mediated decay | - | A0A1B0GT69 | CDS 5' incomplete TSL:5 |
| Col4a1-204 | ENSMUST00000208386.1 | 416 | No protein | Processed transcript | - | - | TSL:3 |
| Col4a1-202 | ENSMUST00000130488.2 | 676 | No protein | Retained intron | - | - | TSL:5 |
| Col4a1-205 | ENSMUST00000209000.1 | 604 | No protein | Retained intron | - | - | TSL:NA |

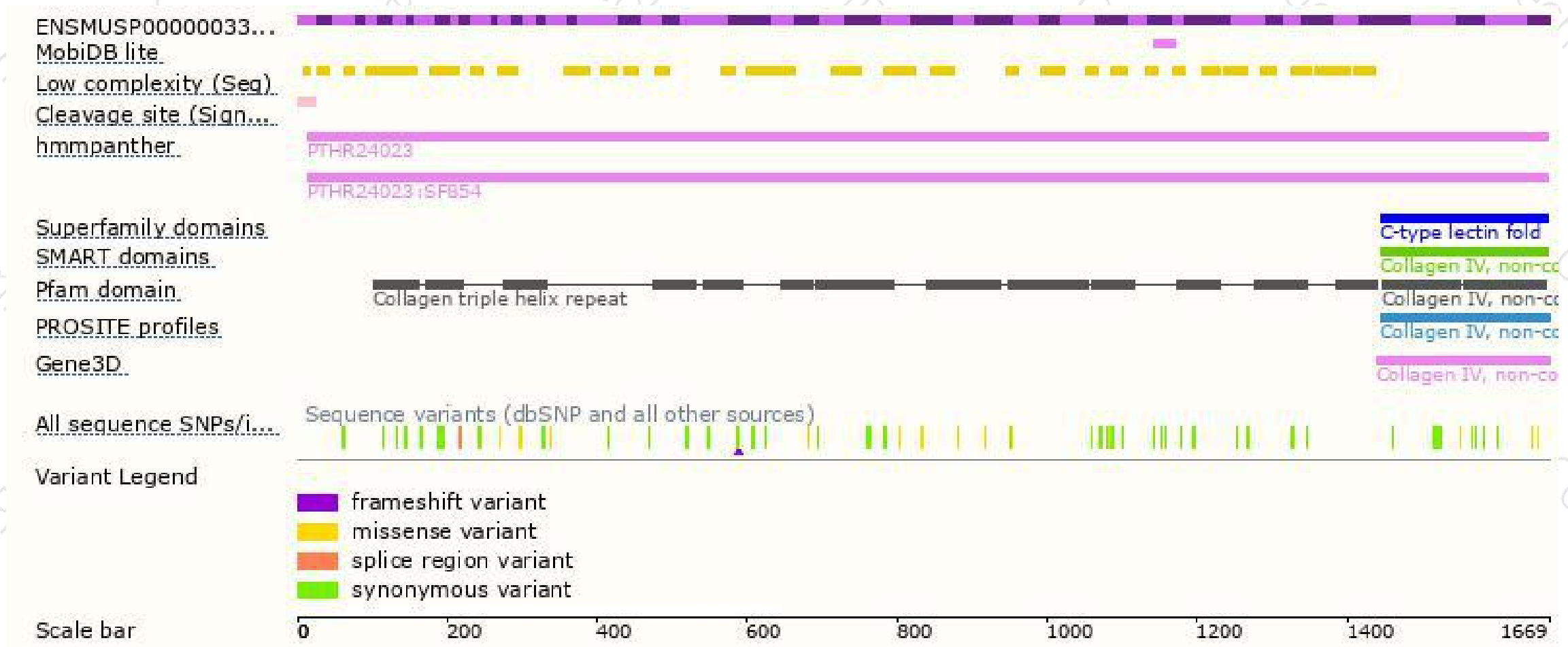
The strategy is based on the design of *Col4a1-201* transcript,The transcription is shown below



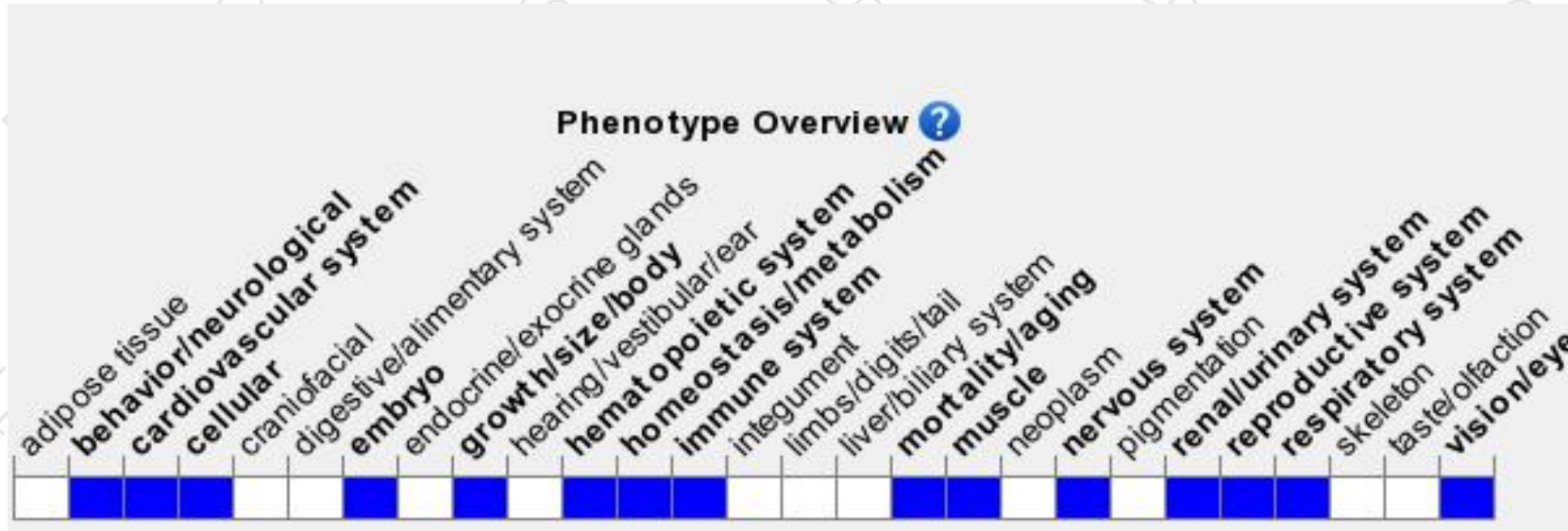
Genomic location distribution



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/>).

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

Tel: 400-9660890

