

***Ccn1(Cyr61)* Cas9-KO Strategy**

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Project Overview

Project Name

Ccn1

Project type

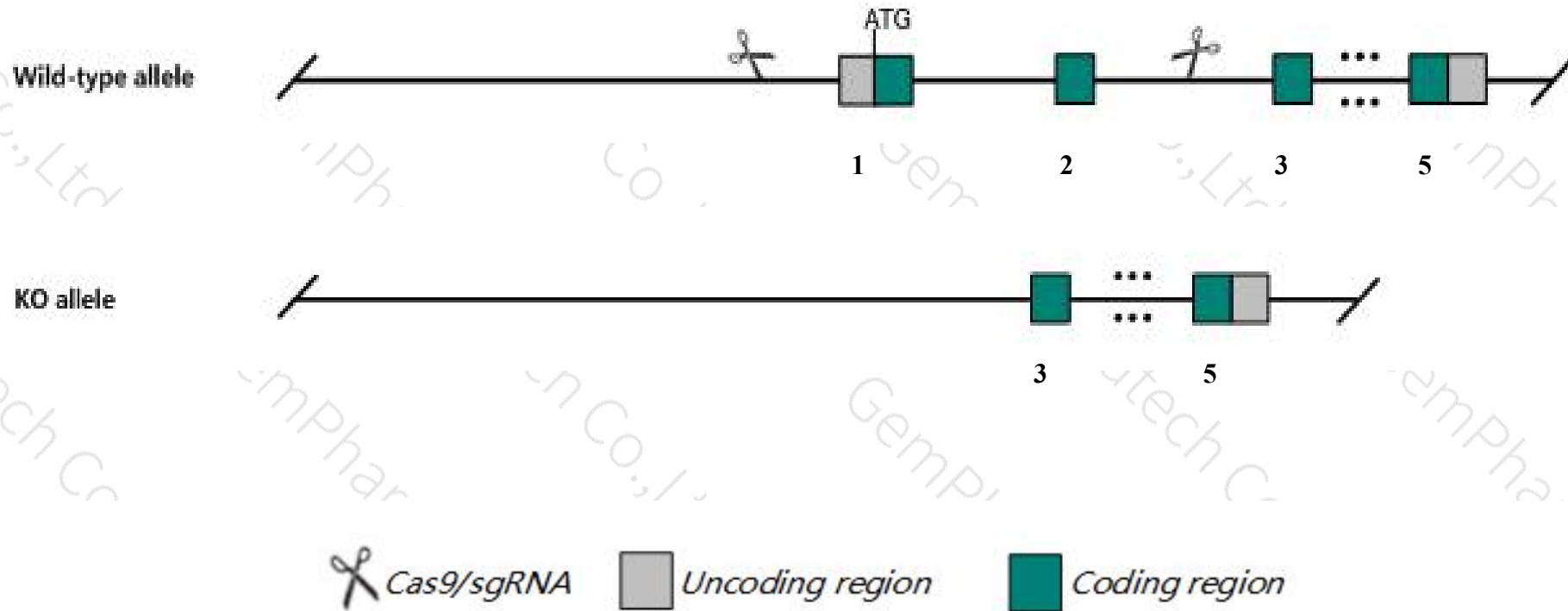
Cas9-KO

Strain background

C57BL/6J

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ccn1* gene. The schematic diagram is as follows:



- The *Ccn1* gene has 1 transcript. According to the structure of *Ccn1* gene, exon1-exon2 of *Ccn1-201* (ENSMUST00000029846.4) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ccn1* 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, Targeted null mice die pre- or perinatally, and none survive beyond 24 hrs of birth. About 30% of embryos die by E10.5 from defects in chorioallantoic fusion, whereas 70% die from placental vascular defects, including impaired allantoic vessel bifurcation, and loss of large-vessel integrity.
- The knockout region contains start codon ATG, there is another ATG that can promote the protein coding.
- The *Ccn1* gene is located on the Chr3. 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 gene transcription and translation processes, all risks cannot be predicted under existing information.

Gene information (NCBI)

Ccn1 cellular communication network factor 1 [Mus musculus (house mouse)]

Gene ID: 16007, updated on 31-Jan-2019

Summary

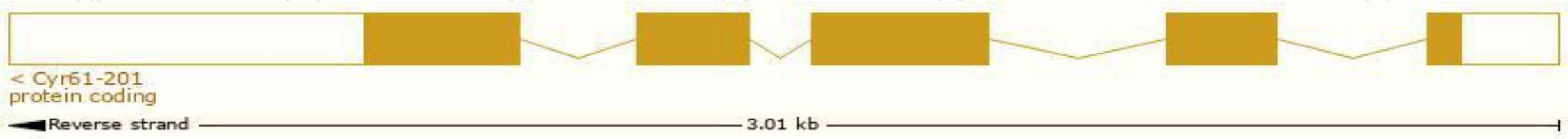
| | |
|--------------------|---|
| Official Symbol | Ccn1 provided by MGI |
| Official Full Name | cellular communication network factor 1 provided by MGI |
| Primary source | MGI:MGI:88613 |
| See related | Ensembl:ENSMUSG00000028195 |
| Gene type | protein coding |
| RefSeq status | VALIDATED |
| Organism | Mus musculus |
| Lineage | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| Also known as | AI325051, Cyr61, Igfbp10 |
| Expression | Broad expression in lung adult (RPKM 41.5), limb E14.5 (RPKM 37.8) and 26 other tissues See more |
| Orthologs | human all |

Transcript information (Ensembl)

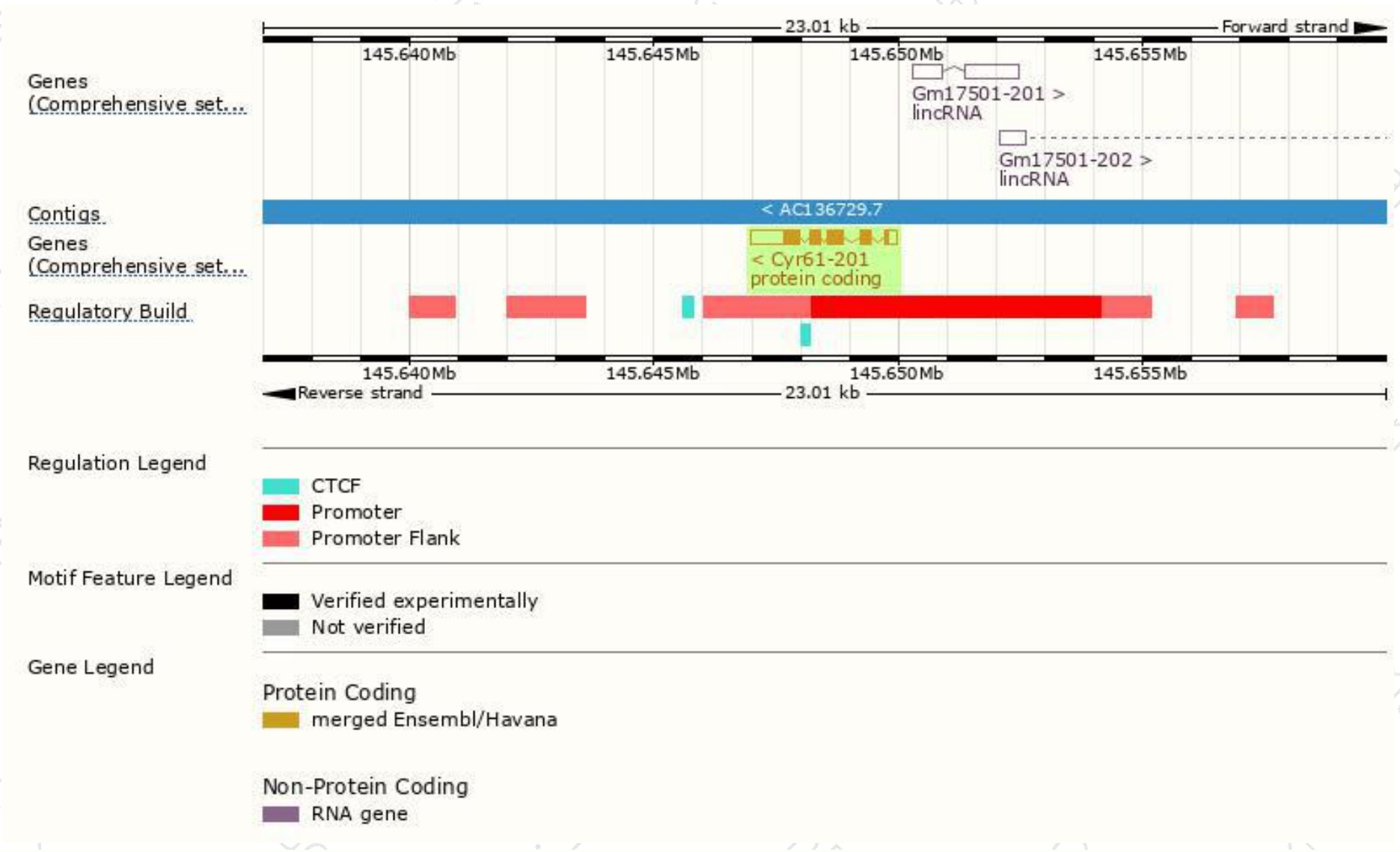
The gene has 1 transcript, and the transcript is shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|----------|--------------------------------------|------|-----------------------|----------------|---------------------------|-------------------------------|-------------------------------|
| Ccn1-201 | ENSMUST00000029846.4 | 2019 | 379aa | Protein coding | CCDS17895 | P18406 Q3TX21 | TSL:1 GENCODE basic APPRIS P1 |

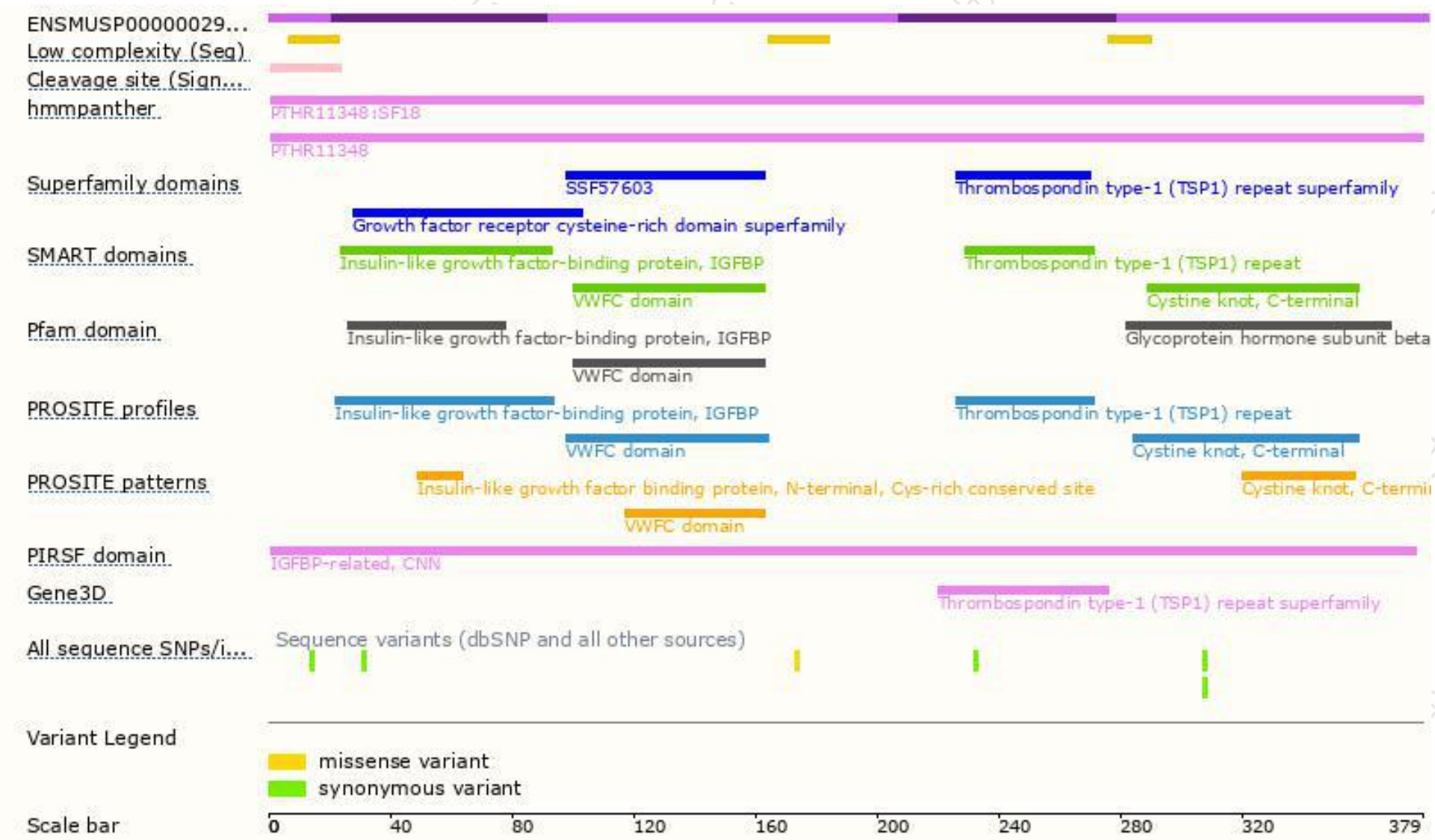
The strategy is based on the design of *Ccn1-201* transcript, The transcription is shown below



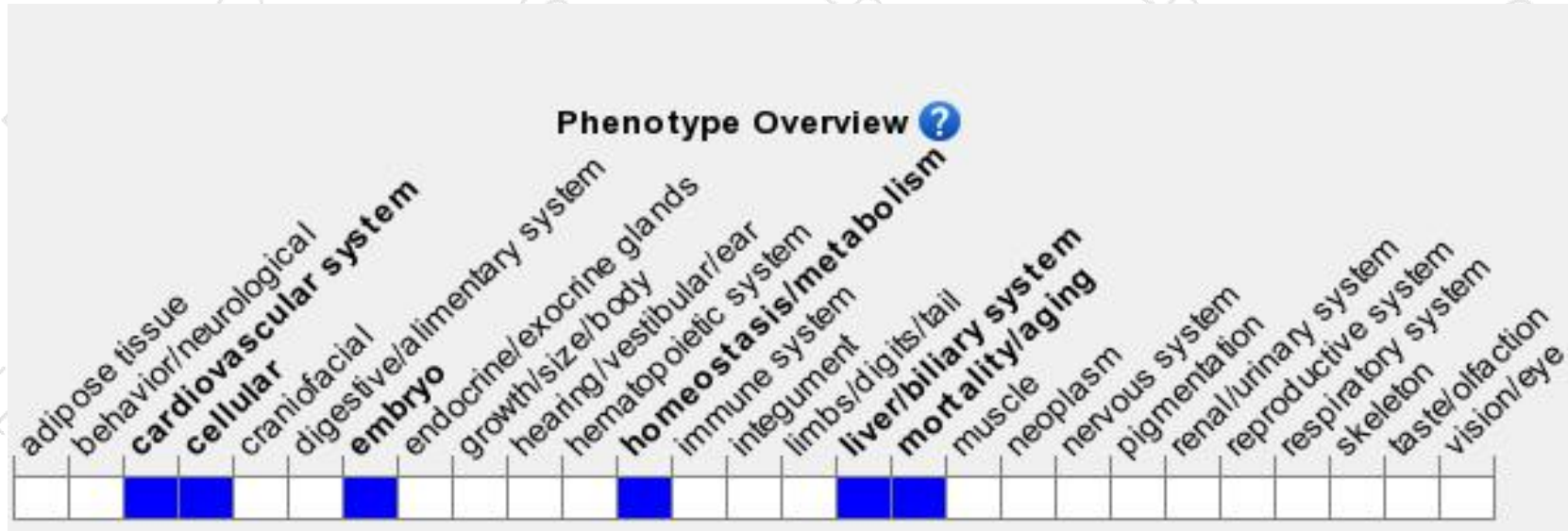
Genomic location distribution



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, Targeted null mice die pre- or perinatally, and none survive beyond 24 hrs of birth.

About 30% of embryos die by E10.5 from defects in chorioallantoic fusion, whereas 70% die from placental vascular defects, including impaired allantoic vessel bifurcation, and loss of large-vessel integrity.

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

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