

***Matn1* Cas9-KO Strategy**

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Project Overview

Project Name

Matn1

Project type

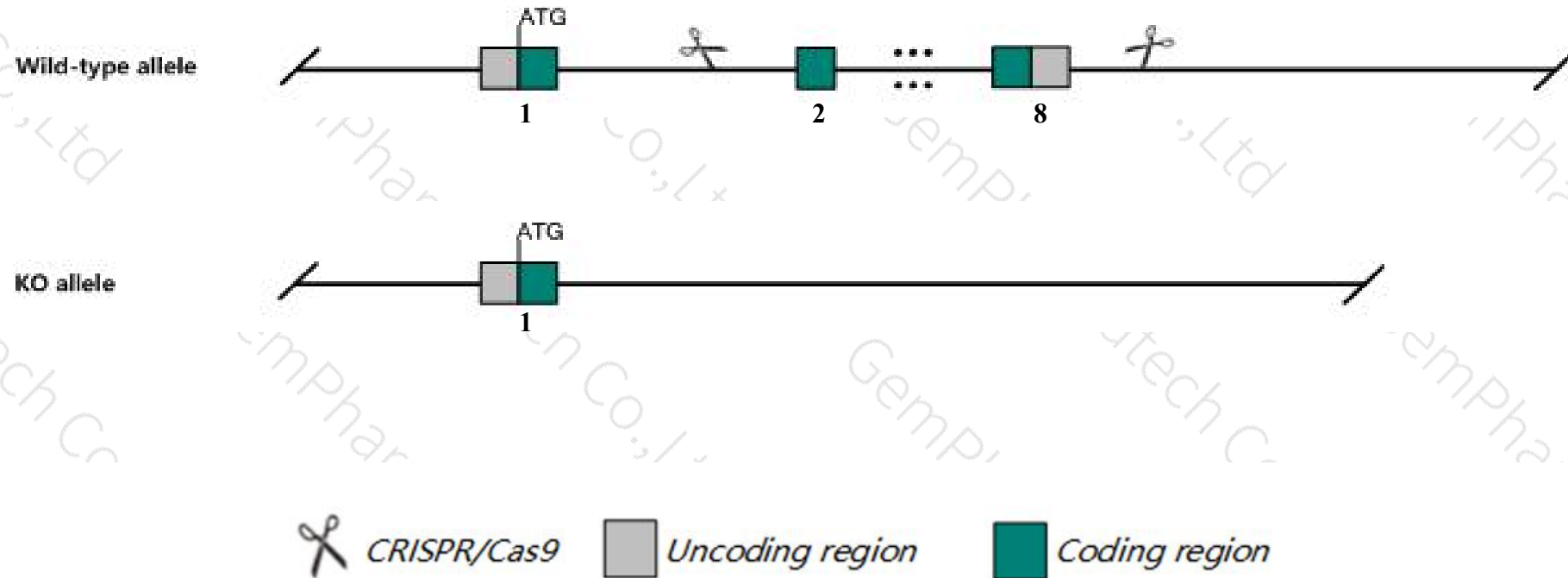
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Matn1* gene. The schematic diagram is as follows:



Technical routes

- The *Matn1* gene has 1 transcript. According to the structure of *Matn1* gene, exon2-exon8 of *Matn1-201* (ENSMUST00000102576.3) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Matn1* gene. The brief process is as follows: CRISPR/Cas9 system were

- According to the existing MGI data, Homozygous null mutants are viable, fertile and display normal cartilage development and endochondral bone formation. Mice homozygous for one targeted allele show alterations in type II collagen fibrillogenesis and fibril organization, in the absence of skeletal defects.
- The *Matn1* gene is located on the Chr4. 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)

Matn1 matrilin 1, cartilage matrix protein [*Mus musculus* (house mouse)]

Gene ID: 17180, updated on 12-Aug-2019

Summary

Official Symbol	Matn1 provided by MGI
Official Full Name	matrilin 1, cartilage matrix protein provided by MGI
Primary source	MGI:MG1:106591
See related	Ensembl:ENSMUSG00000040533
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	CMP; Crtm; Mat1; matrilin-1
Expression	Restricted expression toward limb E14.5 (RPKM 171.3) See more
Orthologs	human all

Genomic context

Location: 4 D2.2; 4 64.09 cM

See Matn1 in [Genome Data Viewer](#)

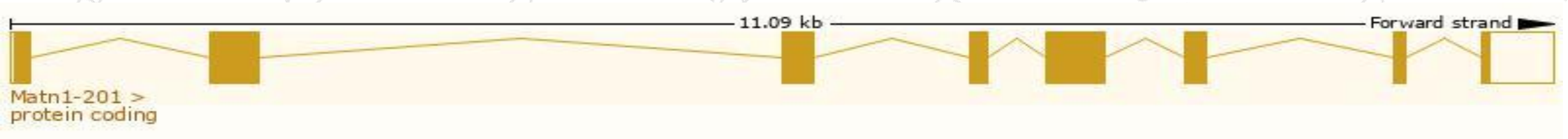
Exon count: 8

Transcript information (Ensembl)

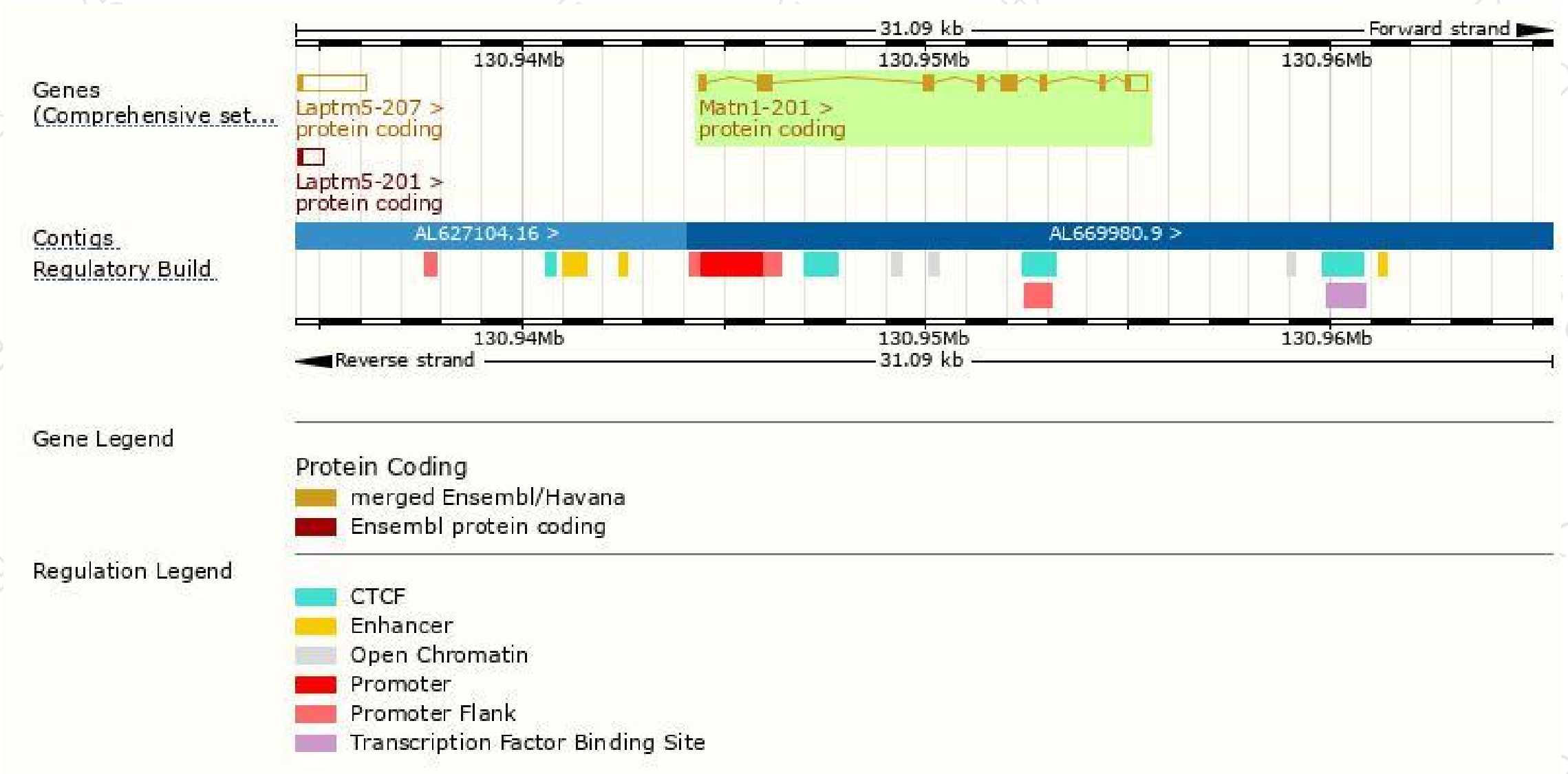
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Matn1-201	ENSMUST00000102576.3	2001	500aa	Protein coding	CCDS18713	P51942	TSL:1 GENCODE basic APPRIS P1

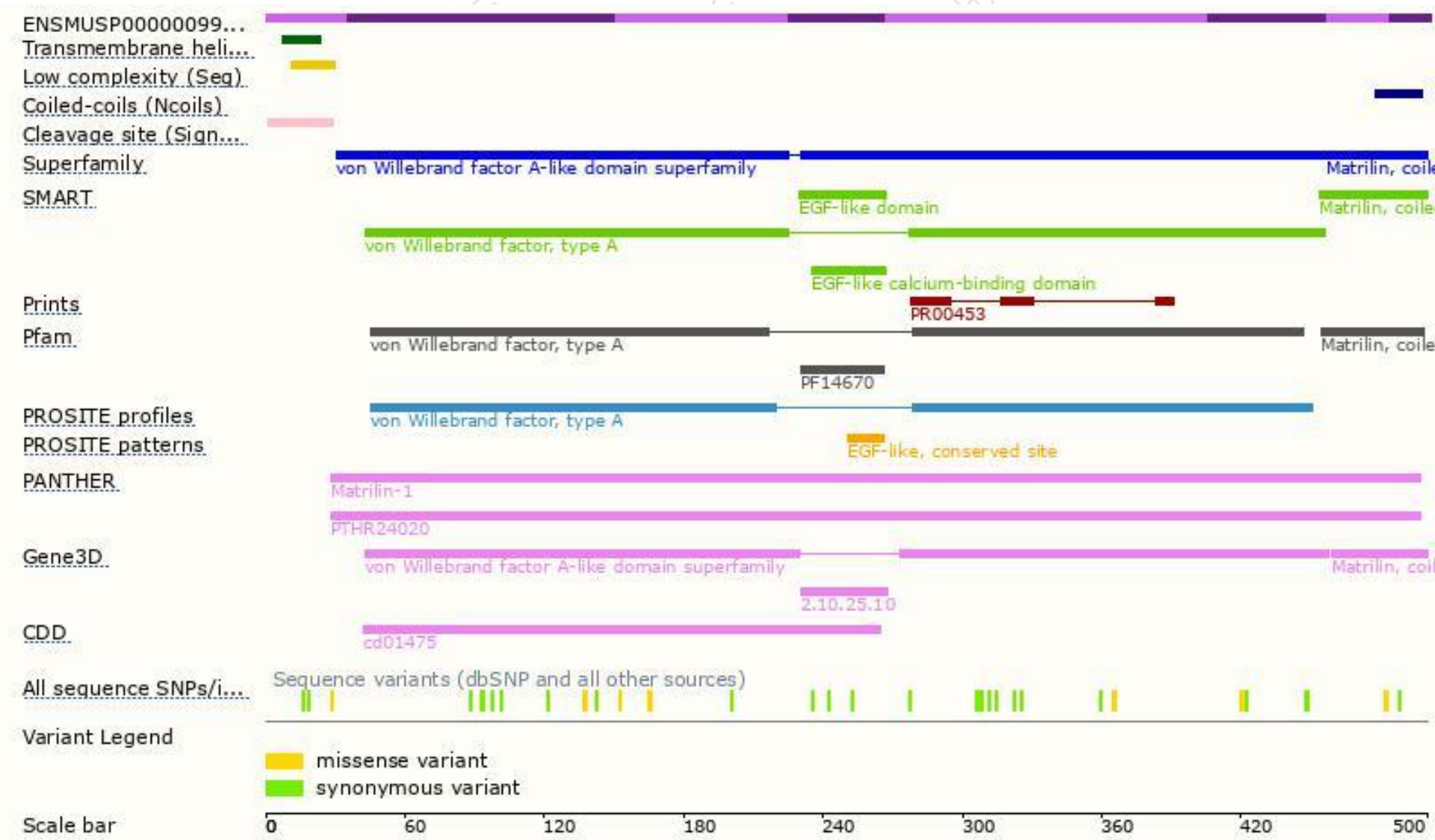
The strategy is based on the design of *Matn1-201* transcript, The transcription is shown below



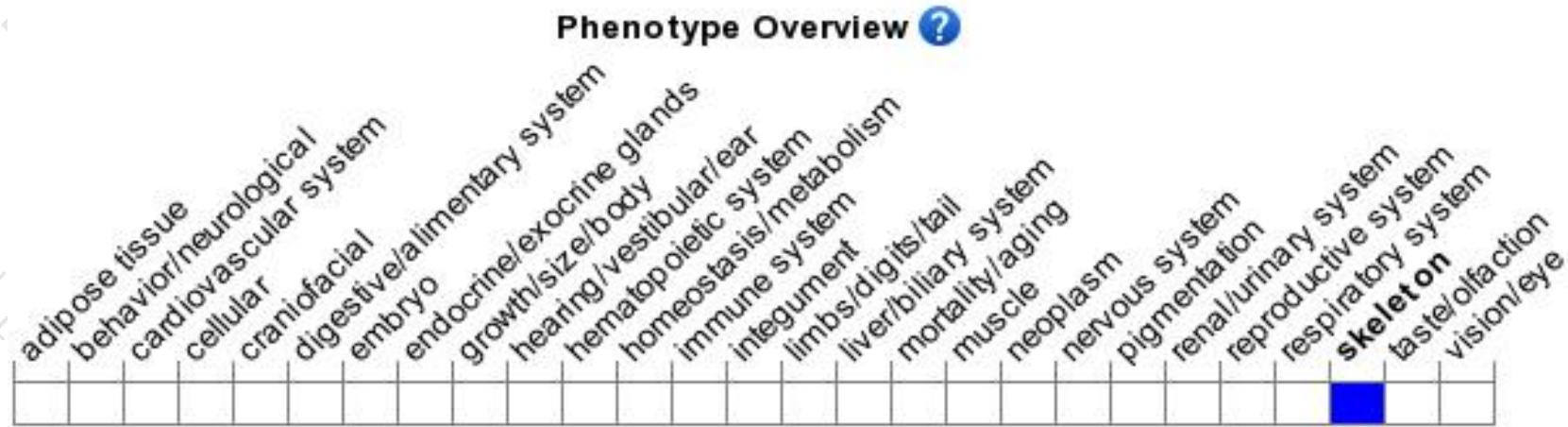
Genomic location distribution



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 are viable, fertile and display normal cartilage development and endochondral bone formation. Mice homozygous for one targeted allele show alterations in type II collagen fibrillogenesis and fibril organization, in the absence of skeletal defects.

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

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