

Zfp36l2 Cas9-CKO Strategy

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Project Overview

Project Name

Zfp36l2

Project type

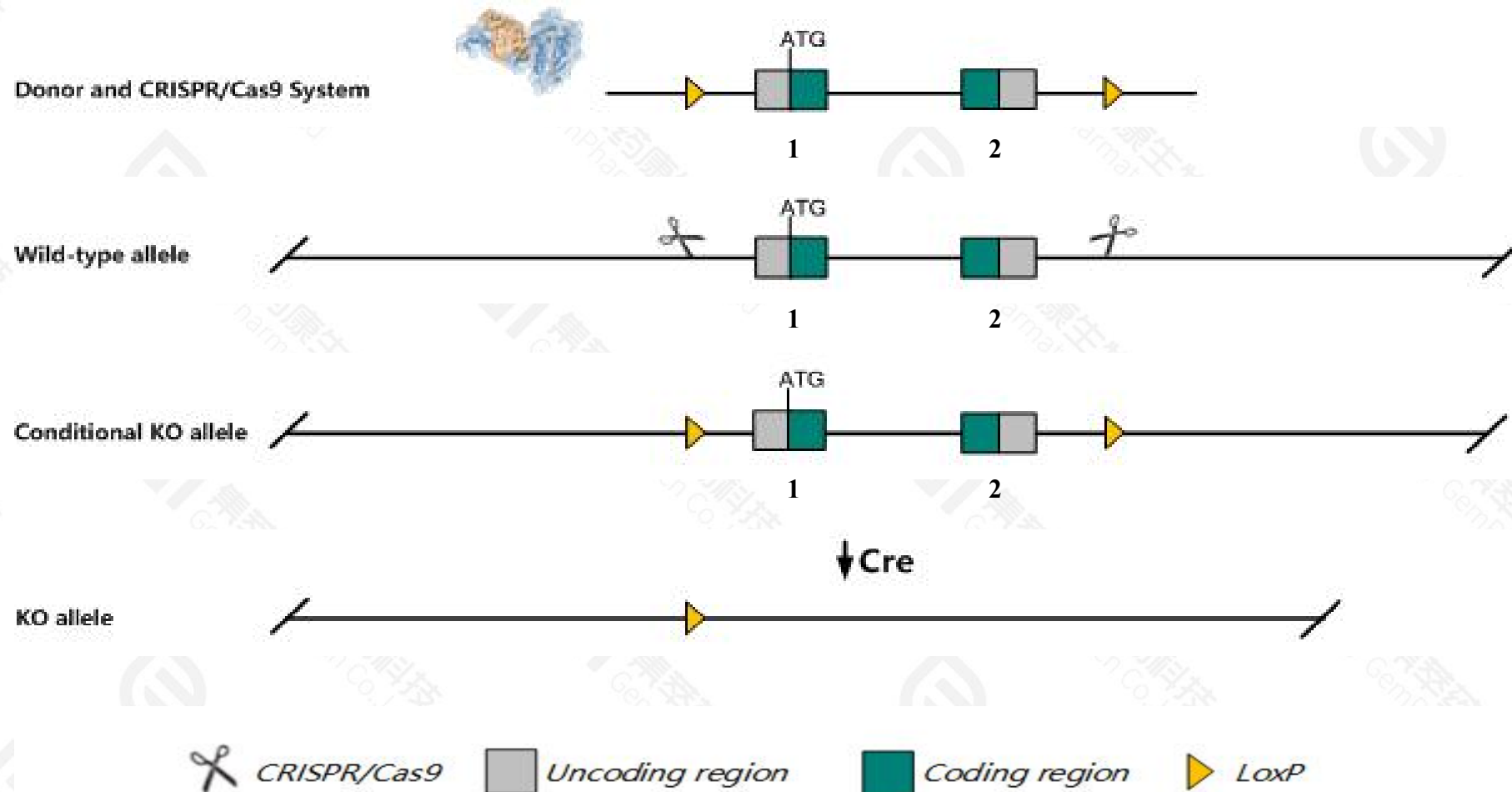
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zfp36l2* gene. The schematic diagram is as follows:



Technical routes

- The *Zfp36l2* gene has 1 transcript. According to the structure of *Zfp36l2* gene, exon1-exon2 of *Zfp36l2*-201(ENSMUST00000060366.7) transcript is recommended as the knockout region. The region contains all 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 *Zfp36l2* 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 expressing decreased levels of an amino-terminal truncated protein display female infertility whereas mice homozygous for a null allele die within two weeks as a result of hematopoietic system defects.
- *Gm36279* gene will be deleted.
- The floxed region is near to the C-terminal of *Thada* gene, this strategy may influence the regulatory function of the C-terminal of *Thada* gene.
- The *Zfp36l2* gene is located on the Chr17. 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)

Zfp36l2 zinc finger protein 36, C3H type-like 2 [Mus musculus (house mouse)]

Gene ID: 12193, updated on 13-Mar-2020

Summary



Official Symbol Zfp36l2 provided by [MGI](#)

Official Full Name zinc finger protein 36, C3H type-like 2 provided by [MGI](#)

Primary source [MGI:MGI:107945](#)

See related [Ensembl:ENSMUSG00000045817](#)

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 Brf2, ERF2, Tis11d

Expression Broad expression in colon adult (RPKM 134.5), ovary adult (RPKM 123.4) and 23 other tissues [See more](#)

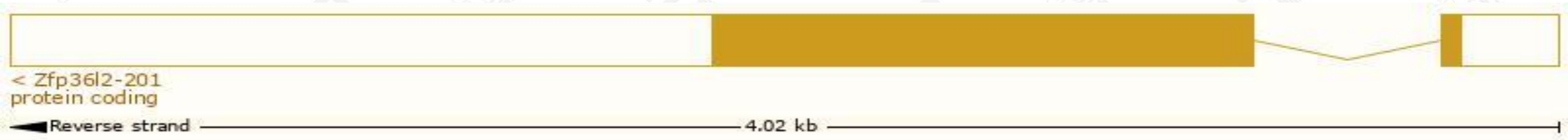
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

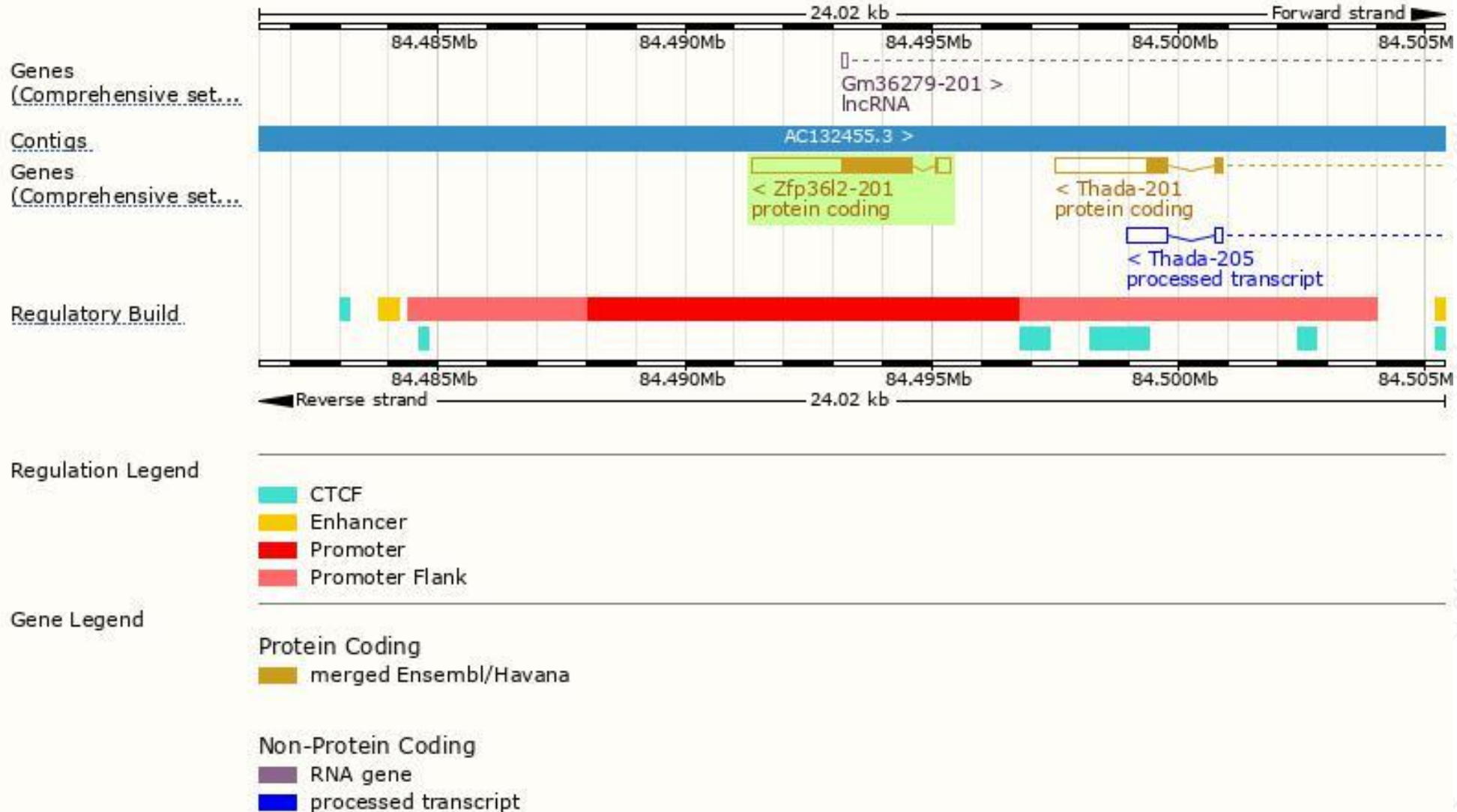
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp36l2-201	ENSMUST00000060366.6	3530	484aa	Protein coding	CCDS28998	P23949	TSL:1 GENCODE basic APPRIS P1

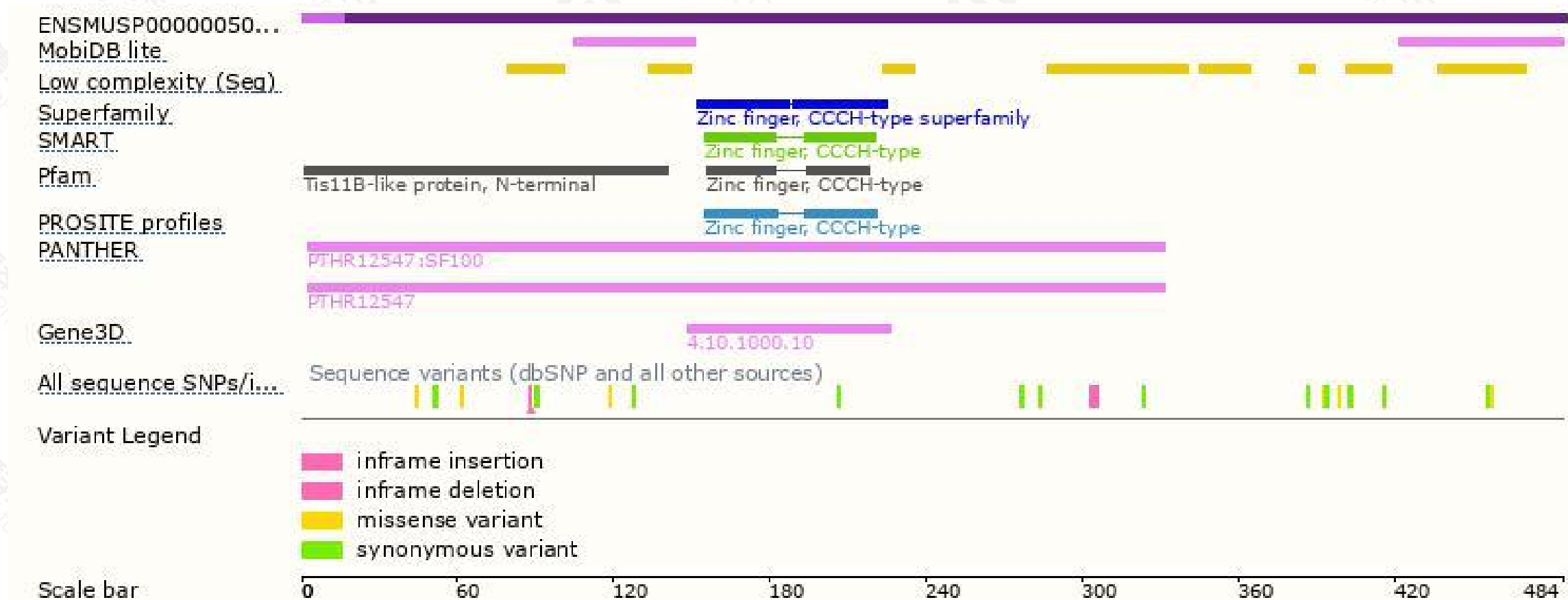
The strategy is based on the design of *Zfp36l2-201* transcript,the transcription is shown below:



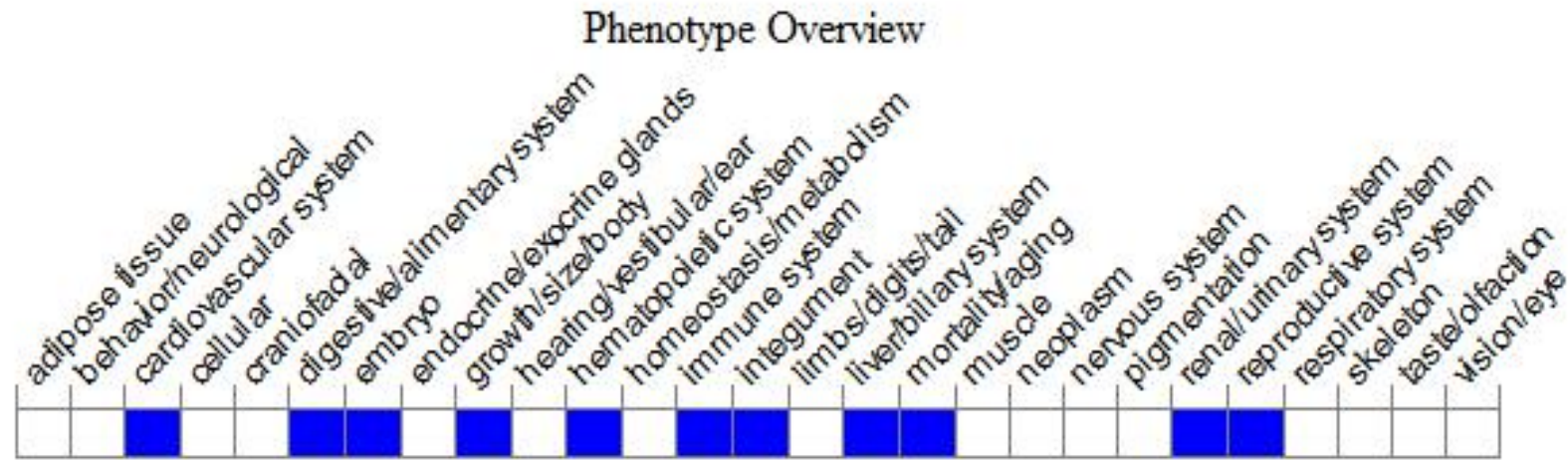
Genomic location distribution



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 expressing decreased levels of an amino-terminal truncated protein display female infertility whereas mice homozygous for a null allele die within two weeks as a result of hematopoietic system defects.

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