

Mak16 Cas9-CKO Strategy

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Design Date: 2021-4-13

Project Overview

Project Name

Mak16

Project type

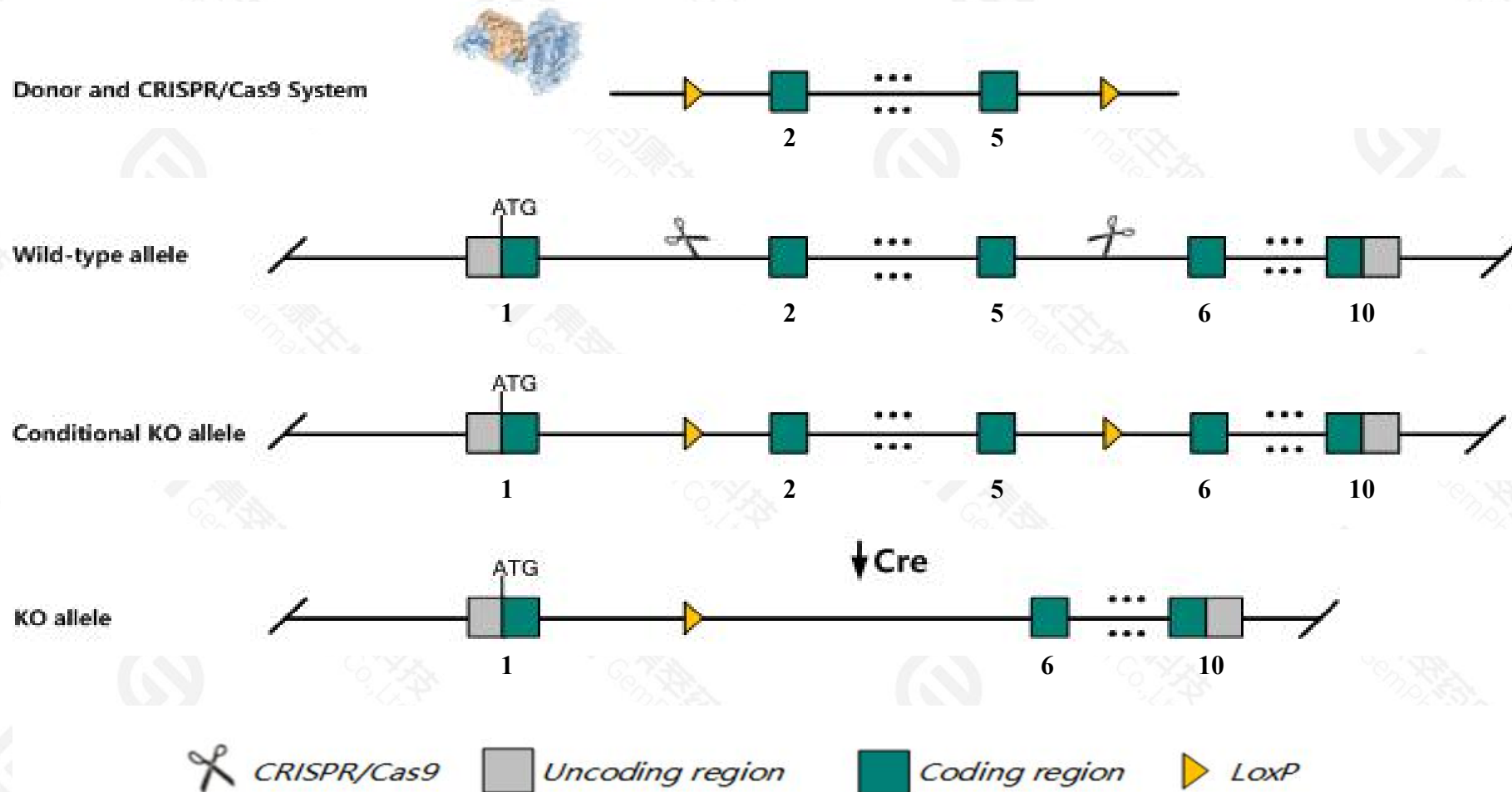
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Mak16* gene. The schematic diagram is as follows:



Technical routes

- The *Mak16* gene has 4 transcripts. According to the structure of *Mak16* gene, exon2-exon5 of *Mak16-201*(ENSMUST00000033983.6) transcript is recommended as the knockout region. The region contains 377bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Mak16* 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.

- The *Mak16* 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)

Mak16 MAK16 homolog [Mus musculus (house mouse)]

Gene ID: 67920, updated on 25-Sep-2020

Summary



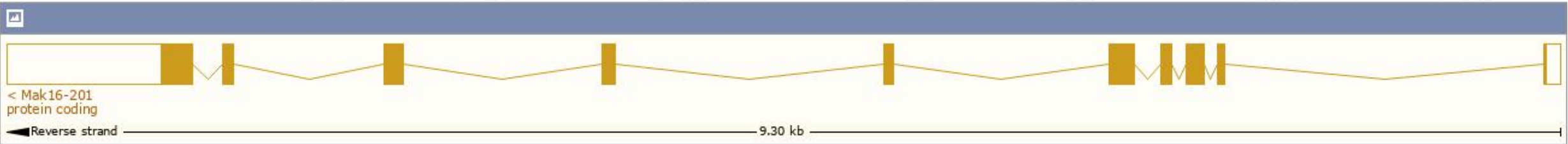
Official Symbol	Mak16 provided by MGI
Official Full Name	MAK16 homolog provided by MGI
Primary source	MGI:MGI:1915170
See related	Ensembl:ENSMUSG00000031578
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2600016B03Rik, AI314911, Rbm1, Rbm13
Expression	Broad expression in CNS E11.5 (RPKM 19.0), liver E14 (RPKM 12.7) and 23 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

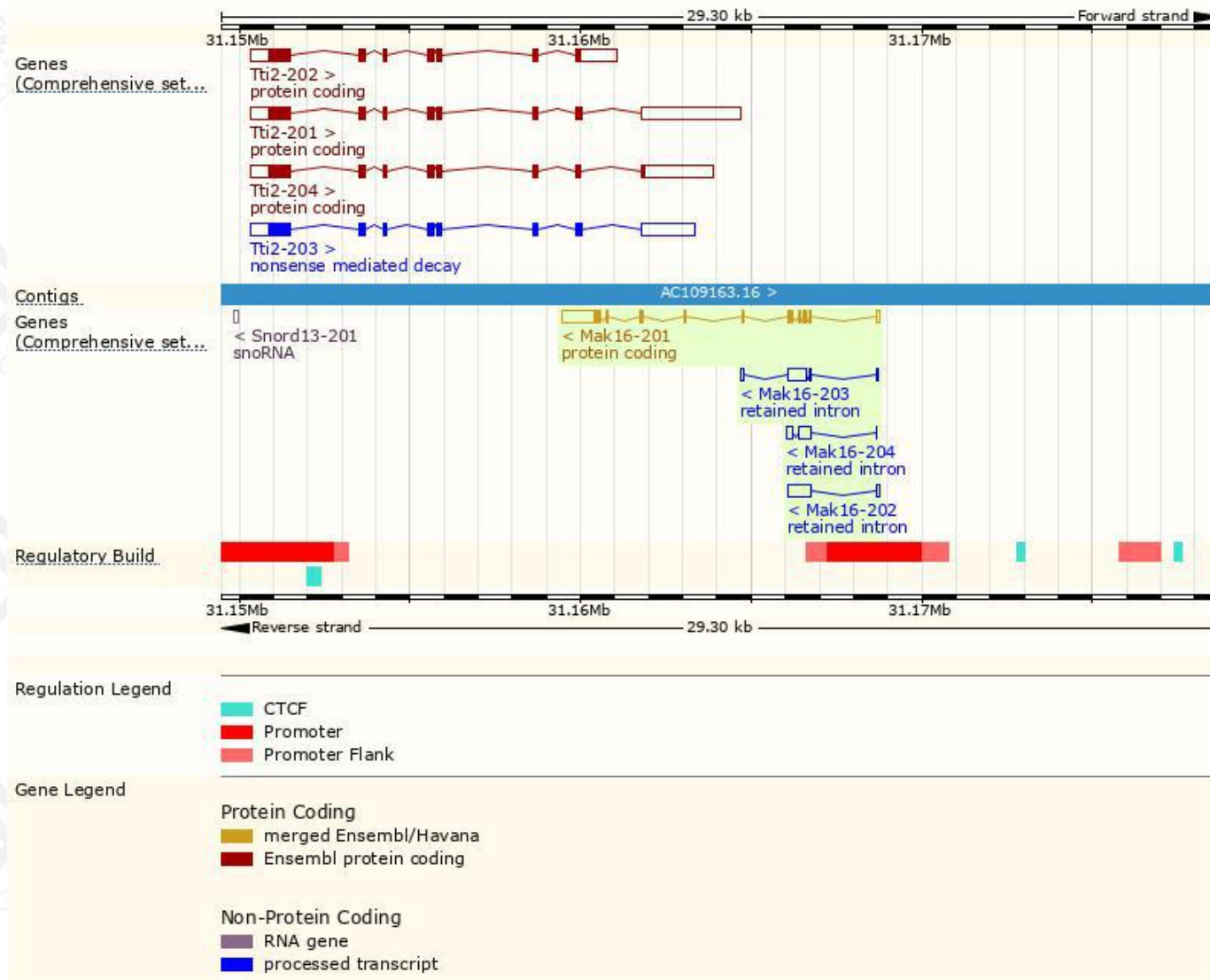
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Mak16-201	ENSMUST00000033983.6	1902	296aa	Protein coding	CCDS22222		TSL:1 , GENCODE basic , APPRIS P1 ,
Mak16-202	ENSMUST00000209991.2	757	No protein	Retained intron	-		TSL:3 ,
Mak16-203	ENSMUST00000210278.2	723	No protein	Retained intron	-		TSL:3 ,
Mak16-204	ENSMUST00000210450.2	575	No protein	Retained intron	-		TSL:3 ,

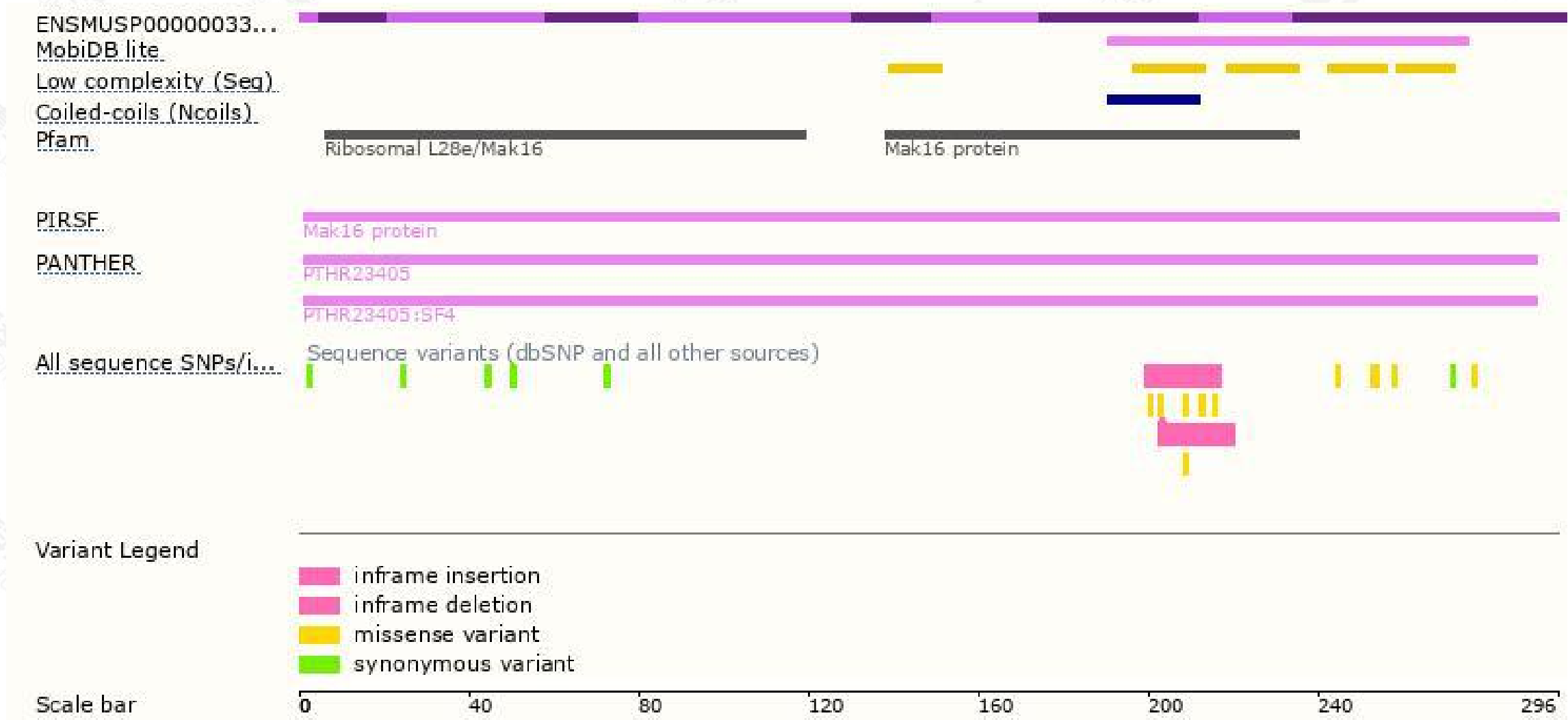
The strategy is based on the design of *Mak16-201* transcript,the transcription is shown below:



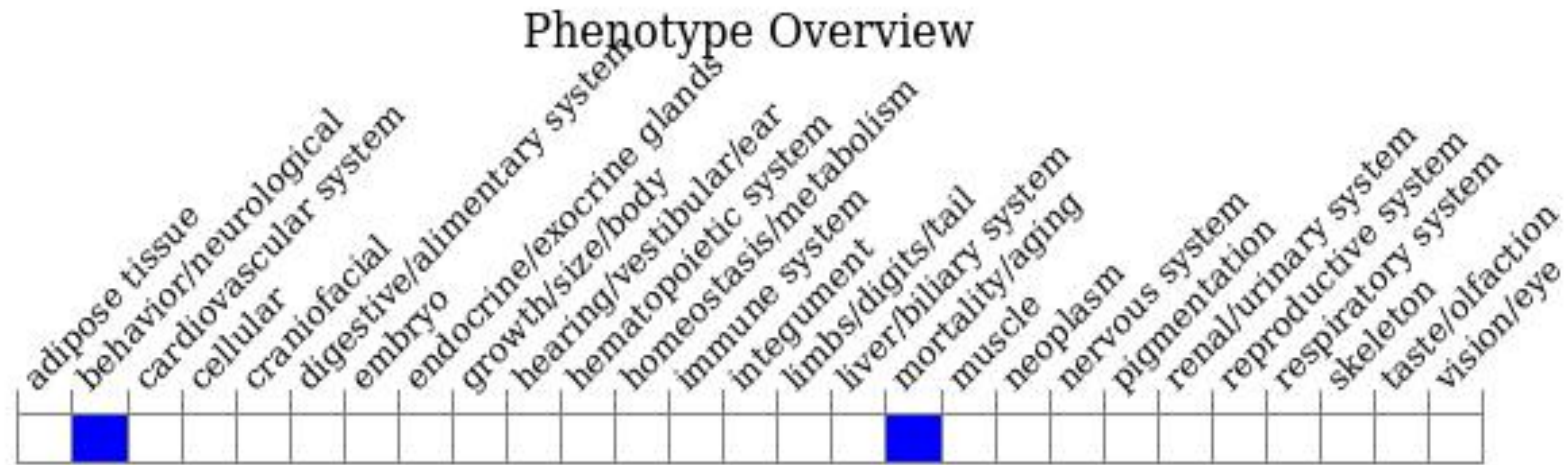
Genomic location distribution



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/>).

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