Serpinf2 Cas9-CKO Strategy

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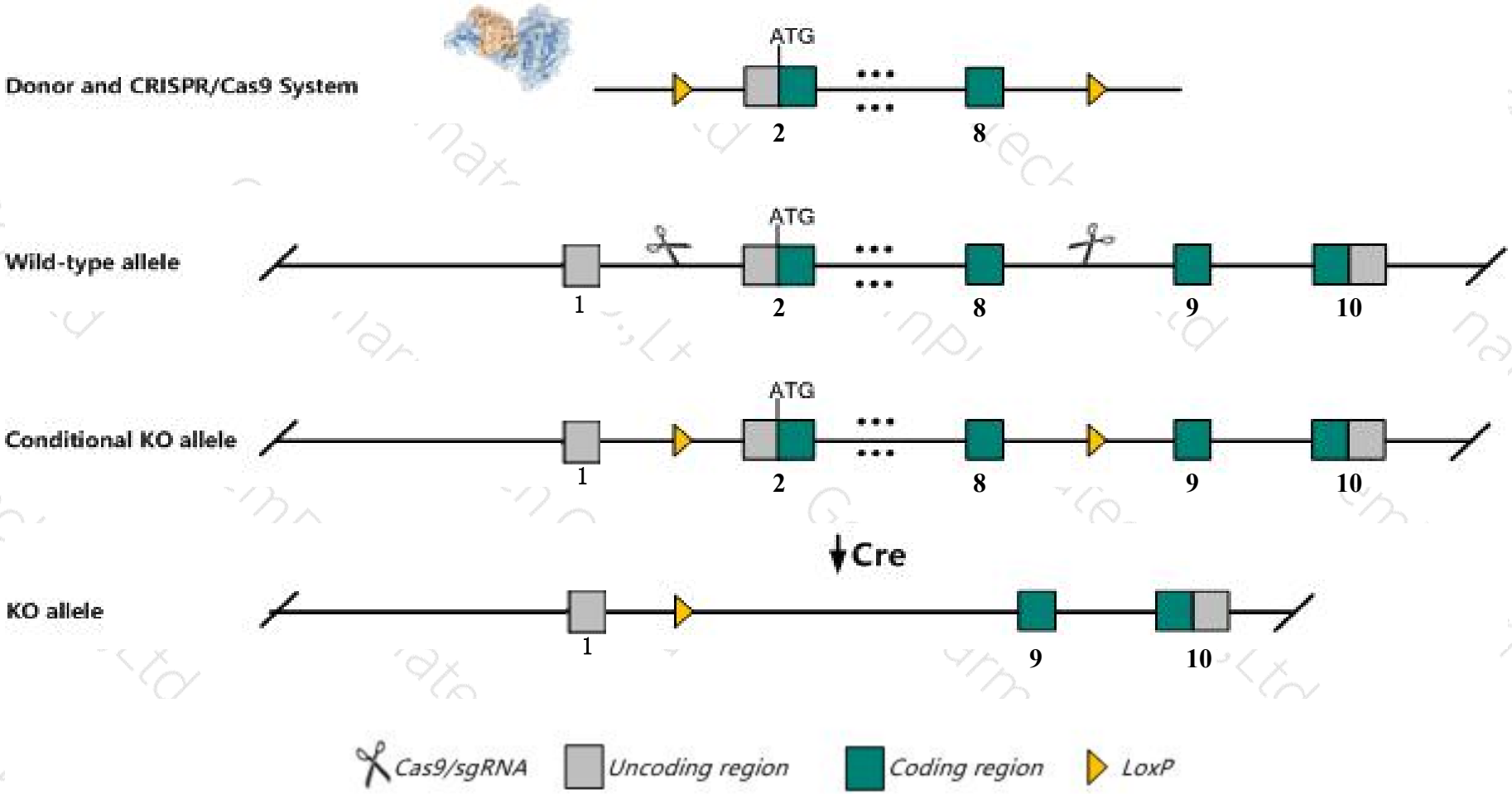
Project Overview

Project Name	***Serpinf2***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Serpinf2* gene. The schematic diagram is as follows:



Technical routes

- The *Serpinf2* gene has 4 transcripts. According to the structure of *Serpinf2* gene, exon2-exon8 of *Serpinf2-201* (ENSMUST00000043696.8) 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 *Serpinf2* 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.

Notice

- According to the existing MGI data,Mice homozygous for disruptions in this gene have an essentially normal phenotype. Spontaneous lysis of blood clots occurs more readily but bleeding times are unaffected.

- The *Serpinf2* gene is located on the Chr11. 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）

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Serpinf2 serine (or cysteine) peptidase inhibitor, clade F, member 2 [*Mus musculus* (house mouse)]

Gene ID: 18816, updated on 1-Feb-2020

Summary

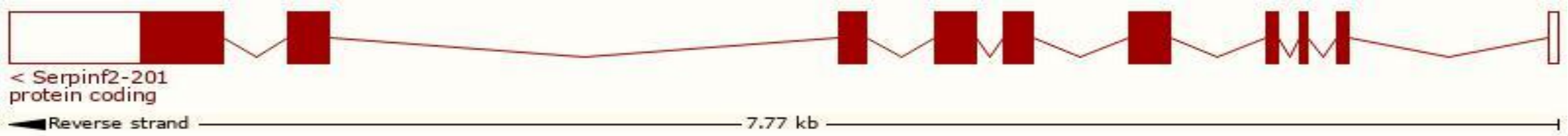
Official Symbol	Serpinf2 provided by MGI
Official Full Name	serine (or cysteine) peptidase inhibitor, clade F, member 2 provided by MGI
Primary source	MGI:MGI:107173
See related	Ensembl:ENSMUSG00000038224
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Pli; AI747498; Serpimf2
Expression	Biased expression in liver adult (RPKM 505.8), liver E18 (RPKM 427.0) and 5 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

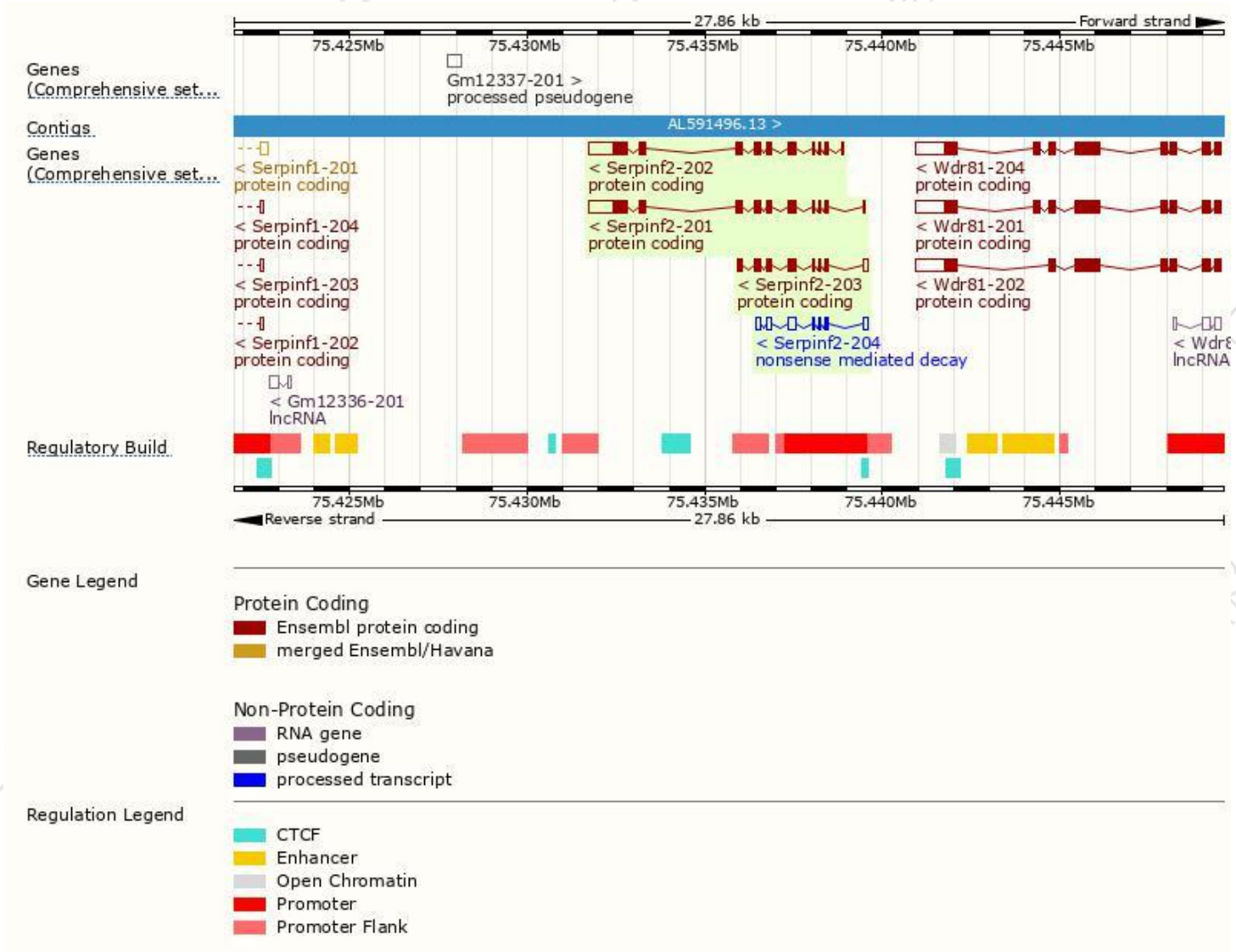
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Serpinf2-201	ENSMUST00000043696.8	2192	491aa	Protein coding	CCDS25046	Q5ND36 Q61247	TSL:1 GENCODE basic APPRIS P1
Serpinf2-202	ENSMUST00000108437.7	2179	491aa	Protein coding	CCDS25046	Q5ND36 Q61247	TSL:1 GENCODE basic APPRIS P1
Serpinf2-203	ENSMUST00000128330.7	981	280aa	Protein coding	-	Q5ND35	CDS 3' incomplete TSL:3
Serpinf2-204	ENSMUST00000142094.1	803	61aa	Nonsense mediated decay	-	E9PXE0	TSL:2

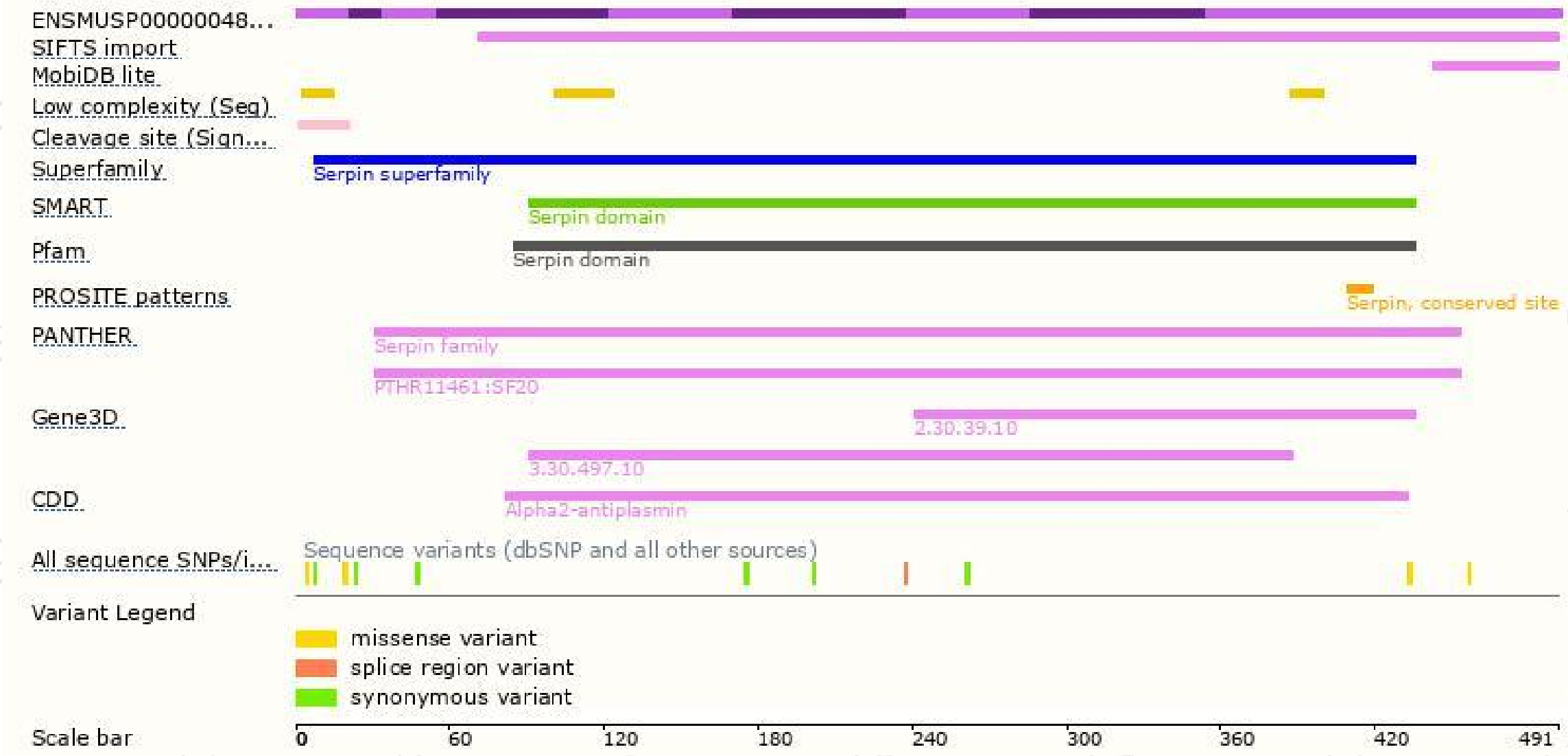
The strategy is based on the design of *Serpinf2-201* transcript,The transcription is shown below



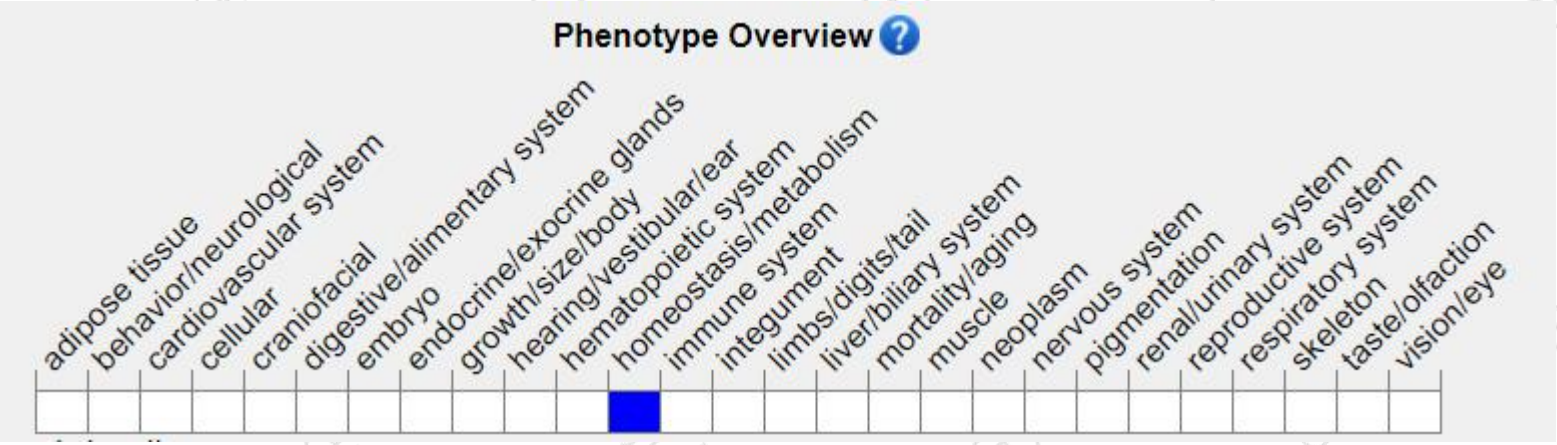
Genomic location distribution



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.

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