Slc22a8 Cas9-CKO Strategy

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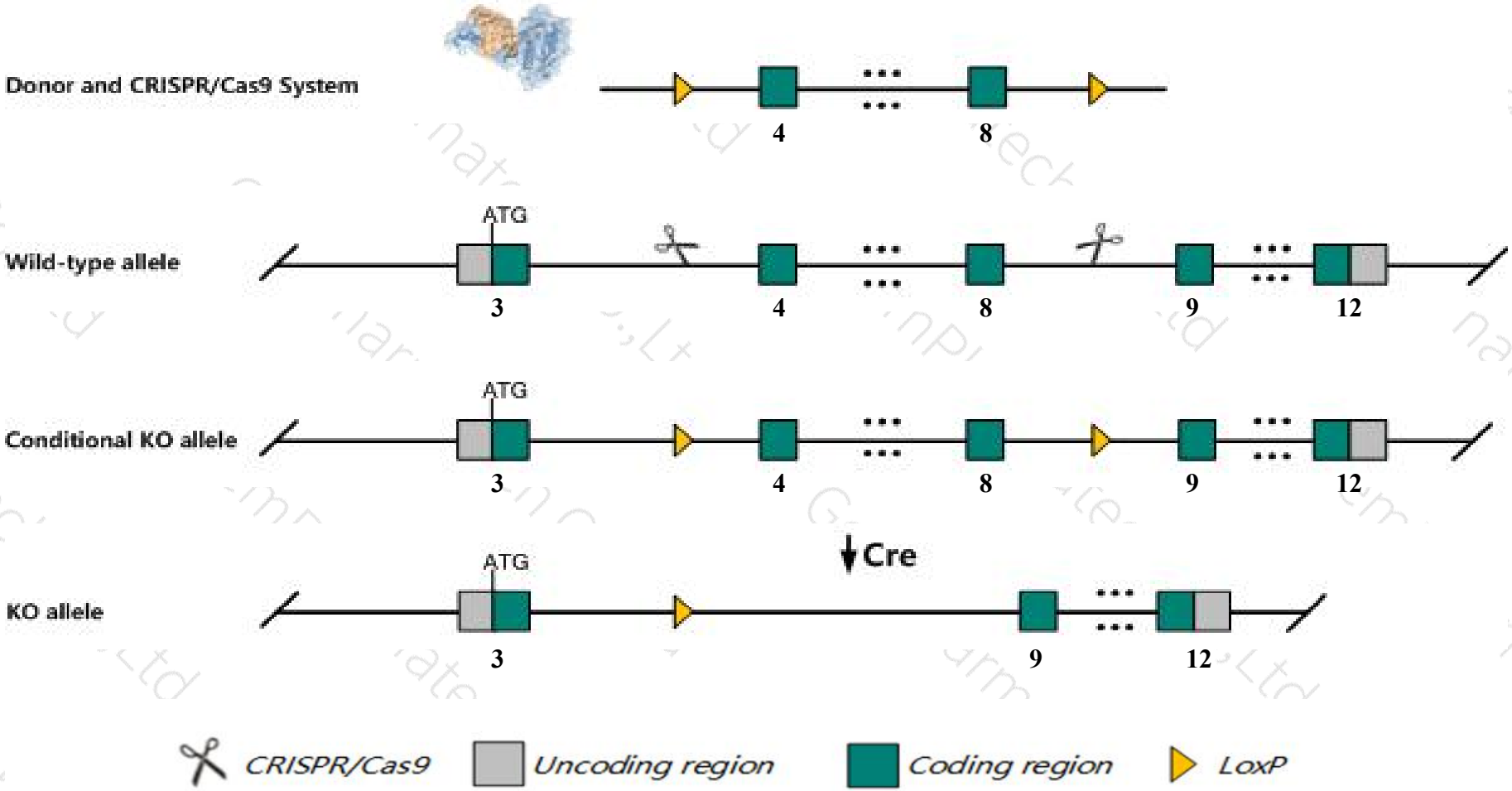
Project Overview

Project Name	***Slc22a8***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc22a8* gene. The schematic diagram is as follows:



Technical routes

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- The *Slc22a8* gene has 3 transcripts. According to the structure of *Slc22a8* gene, exon4-exon8 of *Slc22a8-201* (ENSMUST00000010251.10) transcript is recommended as the knockout region.The region contains 668bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Slc22a8* 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 a null allele exhibit decreased urinary urate levels.

- The *Slc22a8* gene is located on the Chr19. 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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Slc22a8 solute carrier family 22 (organic anion transporter), member 8 [*Mus musculus* (house mouse)]

Gene ID: 19879, updated on 10-Oct-2019

Summary

Official Symbol	Slc22a8 provided by MGI
Official Full Name	solute carrier family 22 (organic anion transporter), member 8 provided by MGI
Primary source	MGI:MGI:1336187
See related	Ensembl:ENSMUSG00000063796
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	Oat3; Roct
Expression	Restricted expression toward kidney adult (RPKM 202.2) See more
Orthologs	human all

Chromosome 19 - NC_000085.6

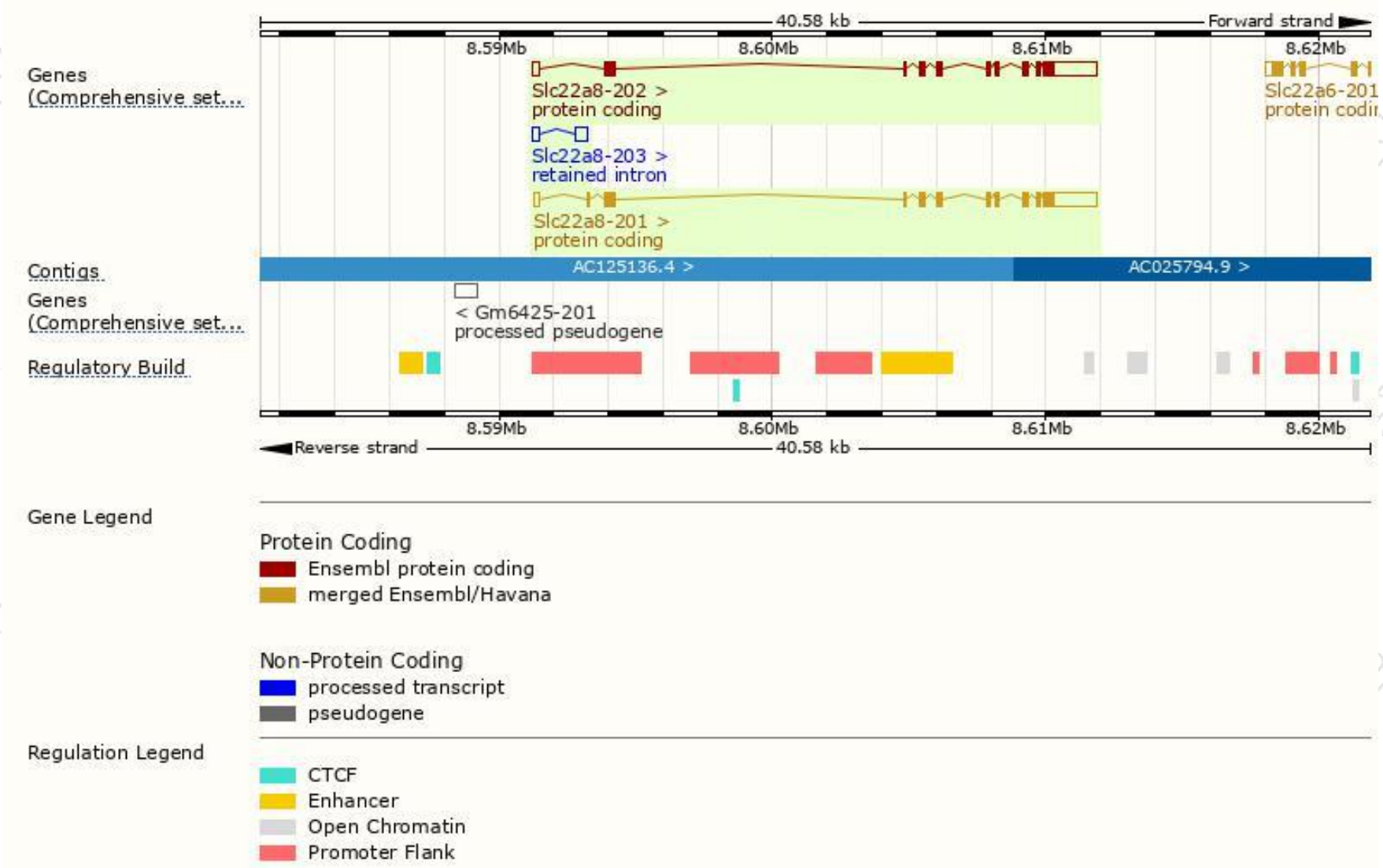
Transcript information（Ensembl）

The gene has 3 transcripts,all transcripts are shown below:

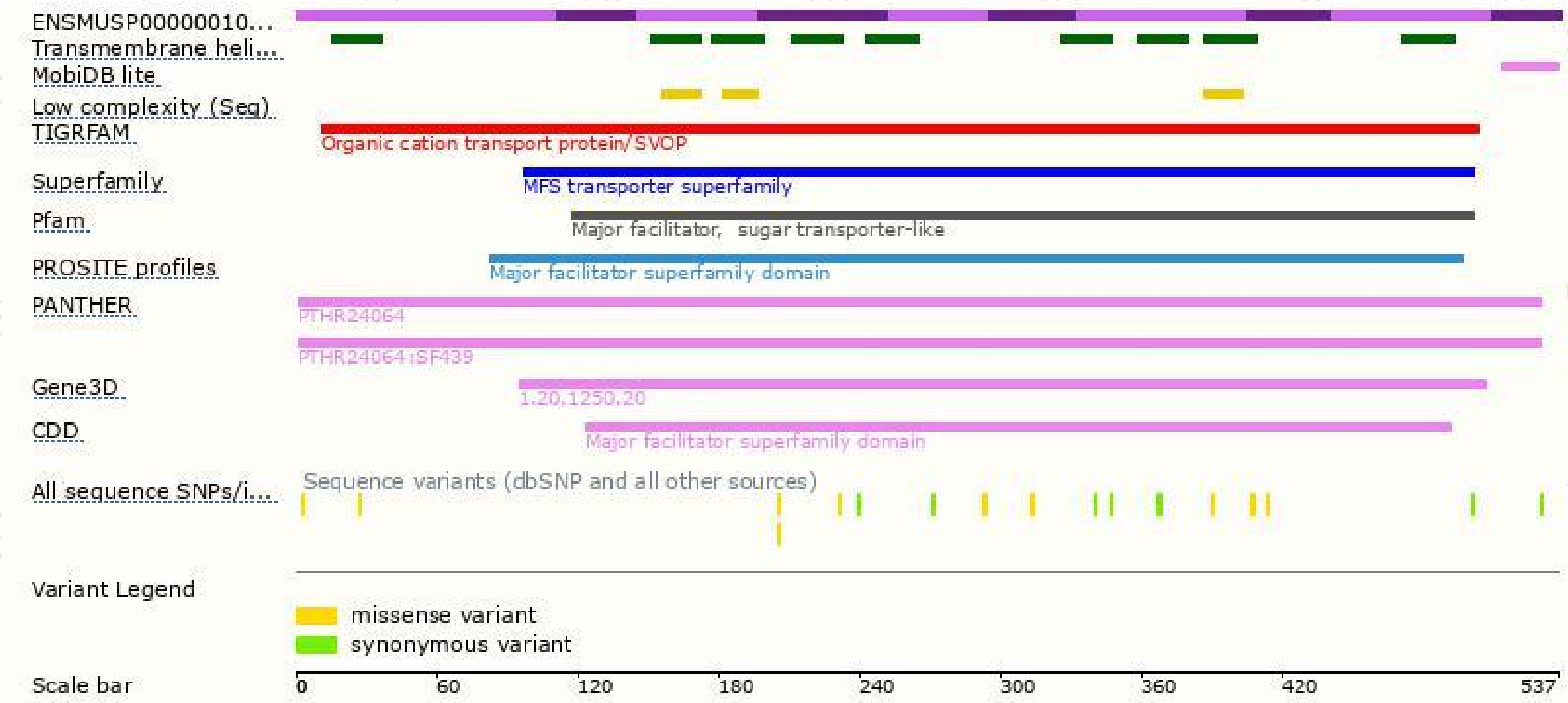
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc22a8-201	ENSMUST00000010251.10	3444	537aa	Protein coding	CCDS29537	O88909	TSL:1 GENCODE basic APPRIS P1
Slc22a8-202	ENSMUST00000170817.1	3398	537aa	Protein coding	CCDS29537	O88909	TSL:5 GENCODE basic APPRIS P1
Slc22a8-203	ENSMUST00000237904.1	664	No protein	Retained intron	-	-	

The strategy is based on the design of *Slc22a8-201* transcript,The transcription is shown below

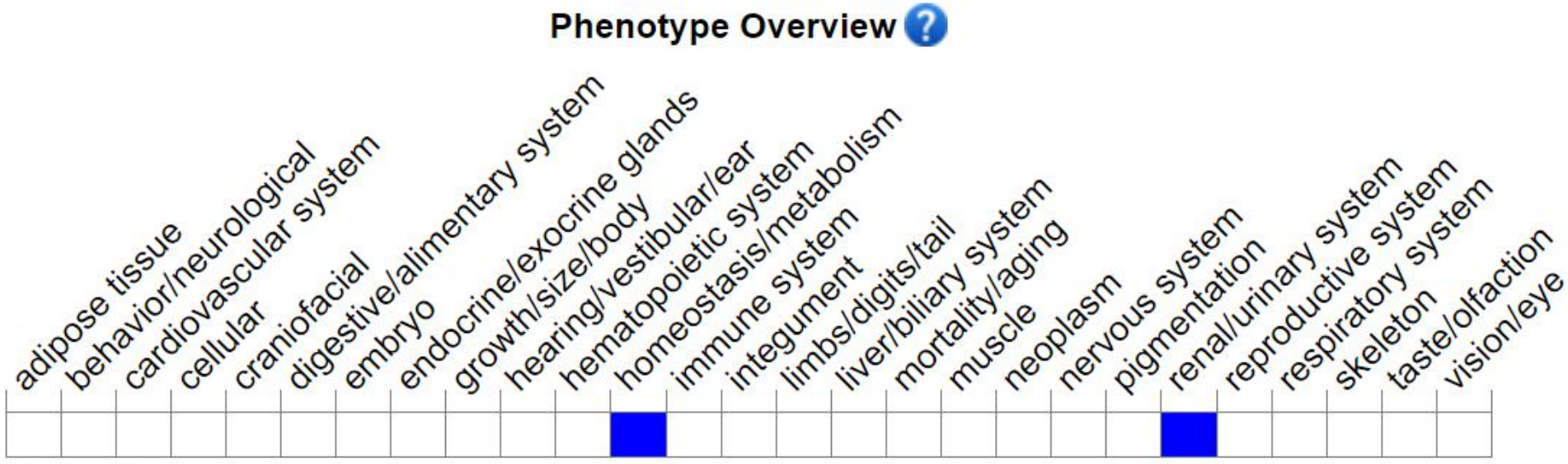
Genomic location distribution



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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