S100a11 Cas9-KO Strategy

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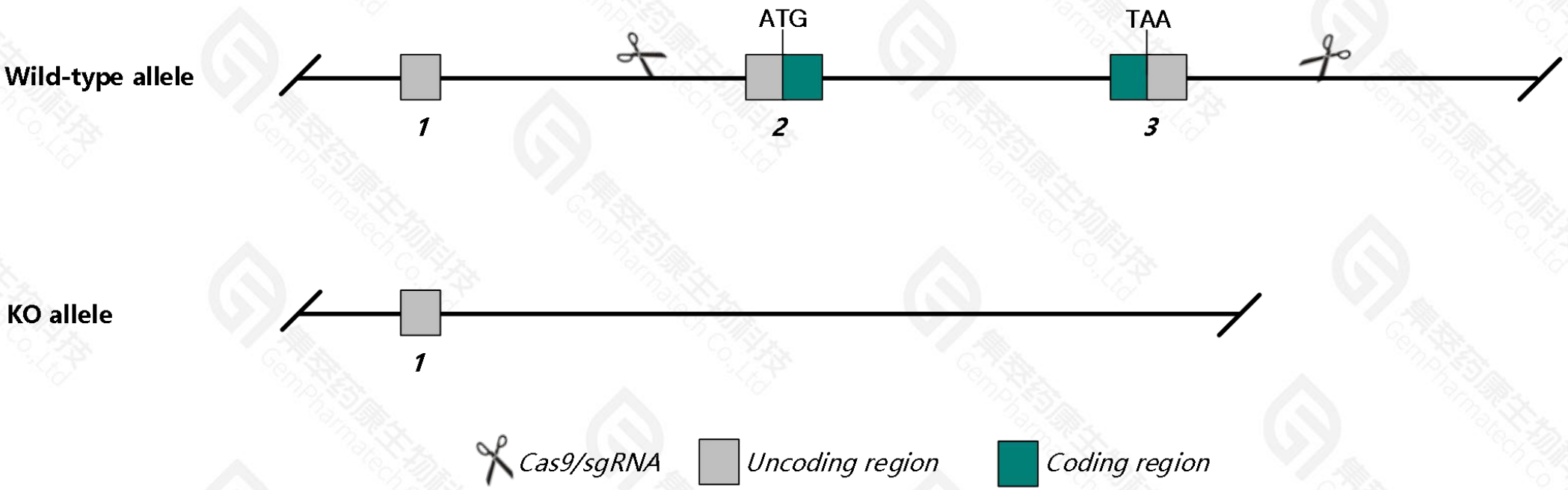
Project Overview

Project Name	***S100a11***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *S100a11* gene. The schematic diagram is as follows:



Technical routes

➢ The *S100a11* gene has 1 transcript. According to the structure of *S100a11* gene, exon2-exon3 of *S100a11-201*(ENSMUST00000029515.5) 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 *S100a11* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice

➢ According to the existing MGI data,mice dwith disruptions in this gene display no obvious phenotype abnormalities other than reduced sperm counts in males.

➢ The *S100a11* gene is located on the Chr3. 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）

S100a11 S100 calcium binding protein A11 [Mus musculus (house mouse)]

Gene ID: 20195, updated on 17-Nov-2020

Summary

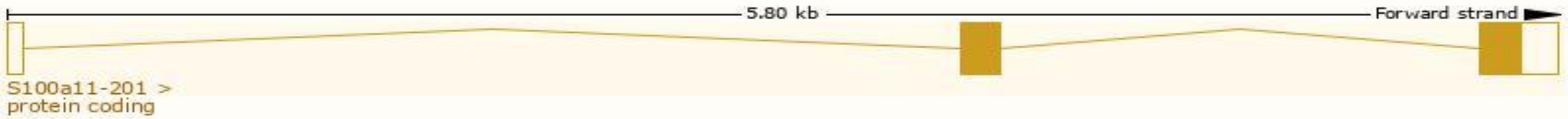
Official Symbol	S100a11 provided by MGI
Official Full Name	S100 calcium binding protein A11 provided byMGI
Primary source	MGI:MGI:1338798
See related	Ensembl:ENSMUSG00000027907
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	EMAPI, Ema, Emap1, S100, S100a, S100a14, S100c, c, cal, calg
Expression	Broad expression in bladder adult (RPKM 206.5), subcutaneous fat pad adult (RPKM 172.6) and 23 other tissuesSee more
Orthologs	human all

Transcript information（Ensembl）

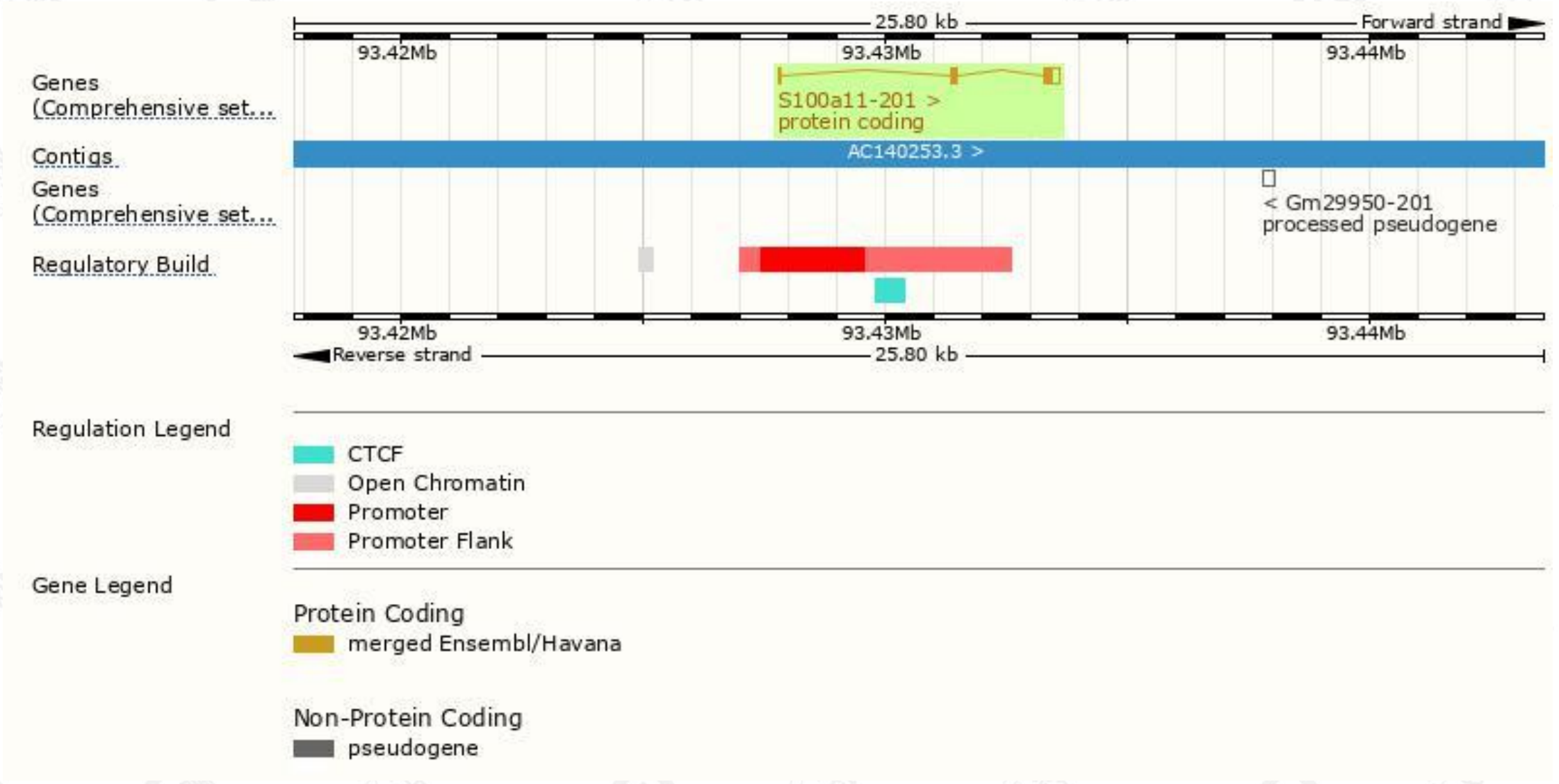
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
S100a11-201	ENSMUST00000029515.5	511	98aa	Protein coding	CCDS38525		TSL:1 , GENCODE basic , APPRIS P1 ,

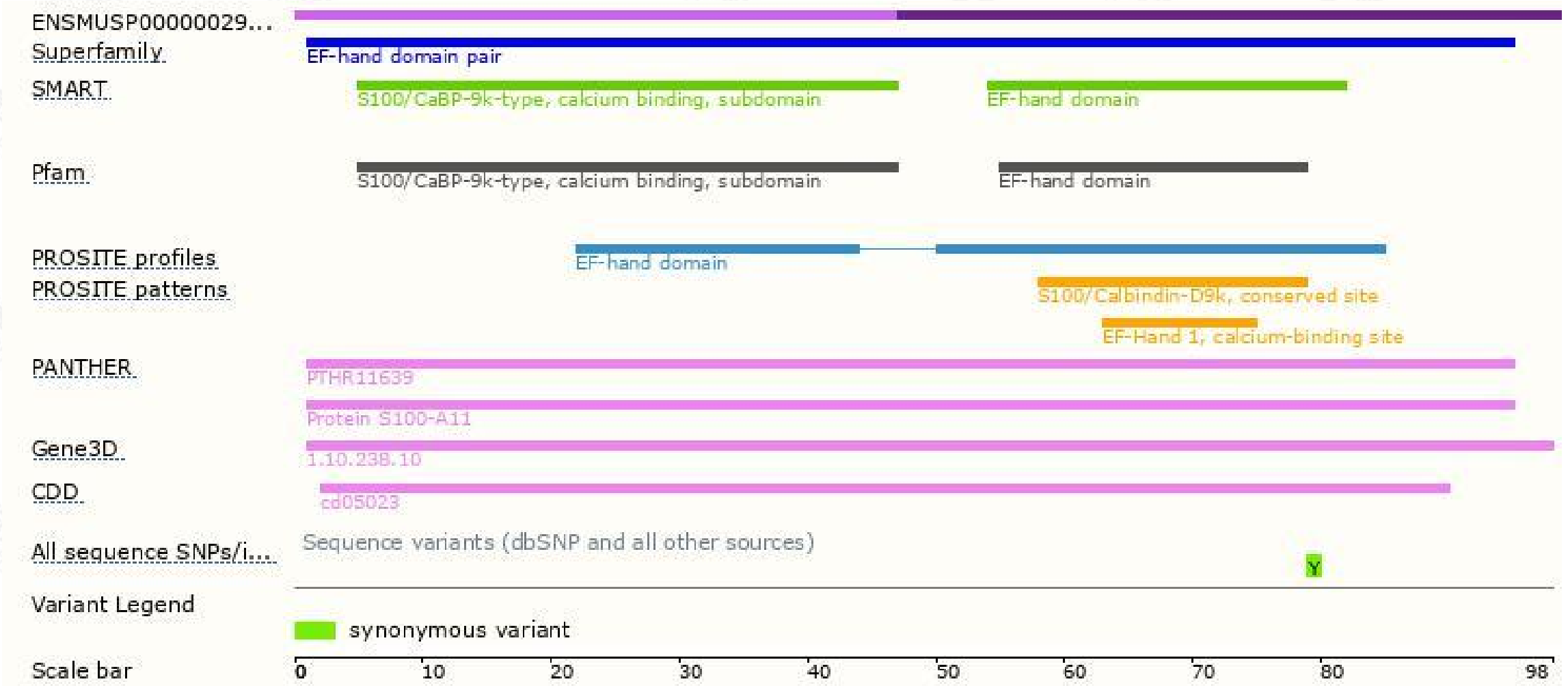
The strategy is based on the design of *S100a11-201* transcript,the transcription is shown below:



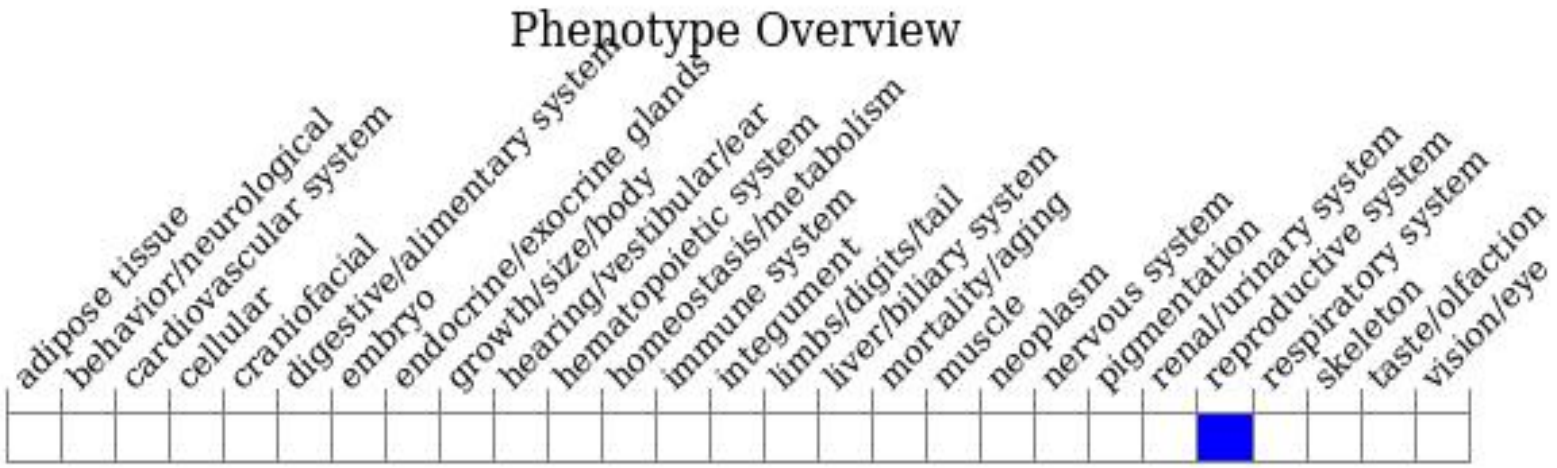
Genomic location distribution



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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