

# *Slco2b1* Cas9-KO Strategy

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# Project Overview

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**Project Name**

*Slco2b1*

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**Project type**

**Cas9-KO**

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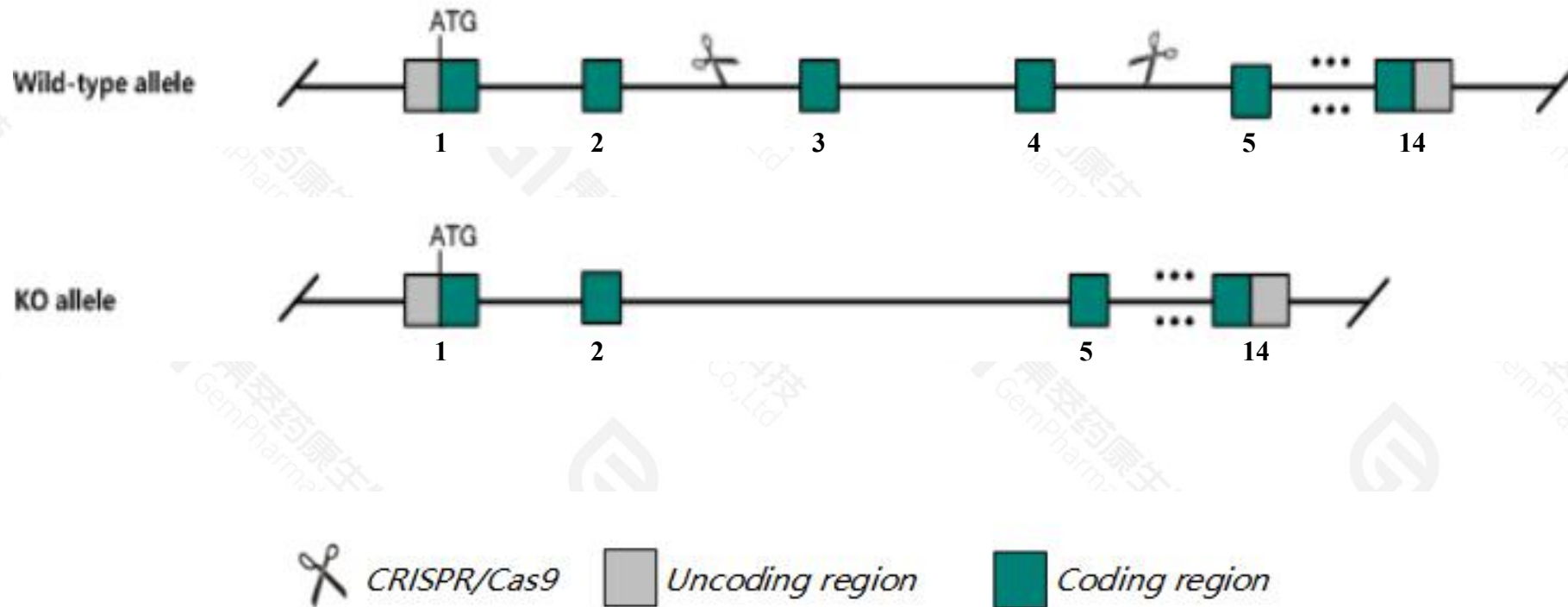
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slco2b1* gene. The schematic diagram is as follows:



# Technical routes

- The *Slco2b1* gene has 9 transcripts. According to the structure of *Slco2b1* gene, exon3-exon4 of *Slco2b1*-203(ENSMUST00000107088.8) transcript is recommended as the knockout region. The region contains 301bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slco2b1* 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.

- The *Slco2b1* gene is located on the Chr7. 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.

## Slco2b1 solute carrier organic anion transporter family, member 2b1 [Mus musculus (house mouse)]

Gene ID: 101488, updated on 12-Jan-2021

### Summary



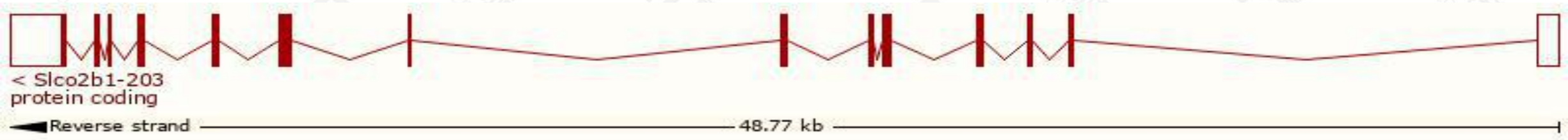
|                           |                                                                                                                                                                           |
|---------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Official Symbol</b>    | Slco2b1 provided by <a href="#">MGI</a>                                                                                                                                   |
| <b>Official Full Name</b> | solute carrier organic anion transporter family, member 2b1 provided by <a href="#">MGI</a>                                                                               |
| <b>Primary source</b>     | <a href="#">MGI:MGI:1351872</a>                                                                                                                                           |
| <b>See related</b>        | <a href="#">Ensembl:ENSMUSG00000030737</a>                                                                                                                                |
| <b>Gene type</b>          | protein coding                                                                                                                                                            |
| <b>RefSeq status</b>      | VALIDATED                                                                                                                                                                 |
| <b>Organism</b>           | <a href="#">Mus musculus</a>                                                                                                                                              |
| <b>Lineage</b>            | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| <b>Also known as</b>      | AI060904, AI852653, OAT, OATP-B, Slc2, Slc21a9                                                                                                                            |
| <b>Expression</b>         | Broad expression in liver adult (RPKM 40.1), mammary gland adult (RPKM 19.5) and 17 other tissues <a href="#">See more</a>                                                |
| <b>Orthologs</b>          | <a href="#">human</a> <a href="#">all</a>                                                                                                                                 |

# Transcript information (Ensembl)

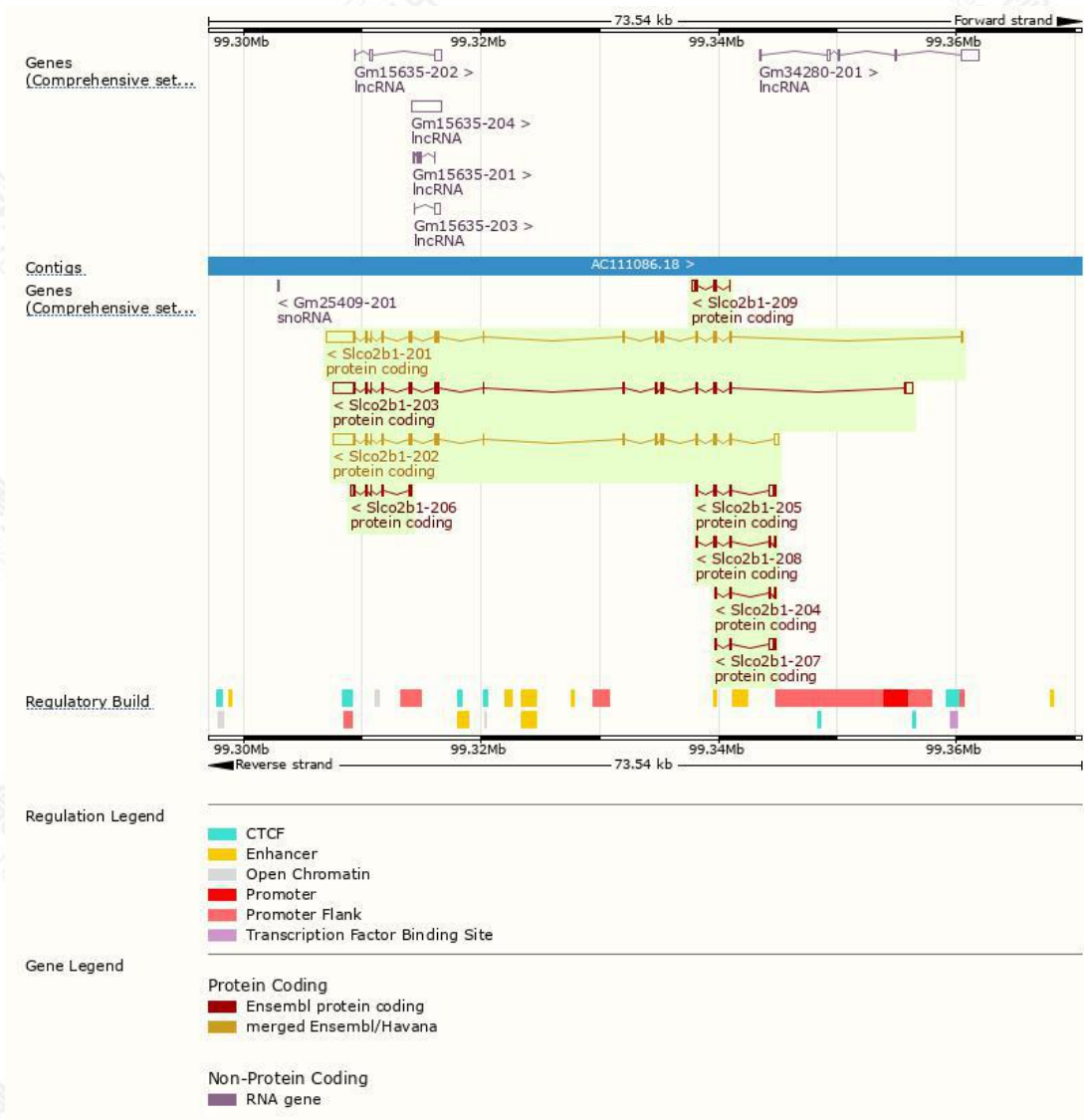
The gene has 9 transcripts,all transcripts are shown below:

| Name        | Transcript ID                         | bp   | Protein               | Biotype        | CCDS                      | UniProt | Flags                                 |
|-------------|---------------------------------------|------|-----------------------|----------------|---------------------------|---------|---------------------------------------|
| Slco2b1-201 | <a href="#">ENSMUST00000032985.11</a> | 4381 | <a href="#">683aa</a> | Protein coding | <a href="#">CCDS21485</a> |         | TSL:1 , GENCODE basic , APPRIS P3 ,   |
| Slco2b1-203 | <a href="#">ENSMUST00000107088.8</a>  | 4369 | <a href="#">693aa</a> | Protein coding | <a href="#">CCDS57567</a> |         | TSL:1 , GENCODE basic , APPRIS ALT2 , |
| Slco2b1-202 | <a href="#">ENSMUST00000107086.9</a>  | 3993 | <a href="#">683aa</a> | Protein coding | <a href="#">CCDS21485</a> |         | TSL:1 , GENCODE basic , APPRIS P3 ,   |
| Slco2b1-205 | <a href="#">ENSMUST00000145381.8</a>  | 796  | <a href="#">150aa</a> | Protein coding | -                         |         | CDS 3' incomplete , TSL:5 ,           |
| Slco2b1-206 | <a href="#">ENSMUST00000154295.2</a>  | 768  | <a href="#">203aa</a> | Protein coding | -                         |         | CDS 5' incomplete , TSL:3 ,           |
| Slco2b1-209 | <a href="#">ENSMUST00000208713.2</a>  | 660  | <a href="#">112aa</a> | Protein coding | -                         |         | CDS 5' incomplete , TSL:5 ,           |
| Slco2b1-208 | <a href="#">ENSMUST00000208225.2</a>  | 636  | <a href="#">150aa</a> | Protein coding | -                         |         | CDS 3' incomplete , TSL:5 ,           |
| Slco2b1-207 | <a href="#">ENSMUST00000207090.2</a>  | 569  | <a href="#">81aa</a>  | Protein coding | -                         |         | CDS 3' incomplete , TSL:3 ,           |
| Slco2b1-204 | <a href="#">ENSMUST00000137914.2</a>  | 405  | <a href="#">85aa</a>  | Protein coding | -                         |         | CDS 3' incomplete , TSL:5 ,           |

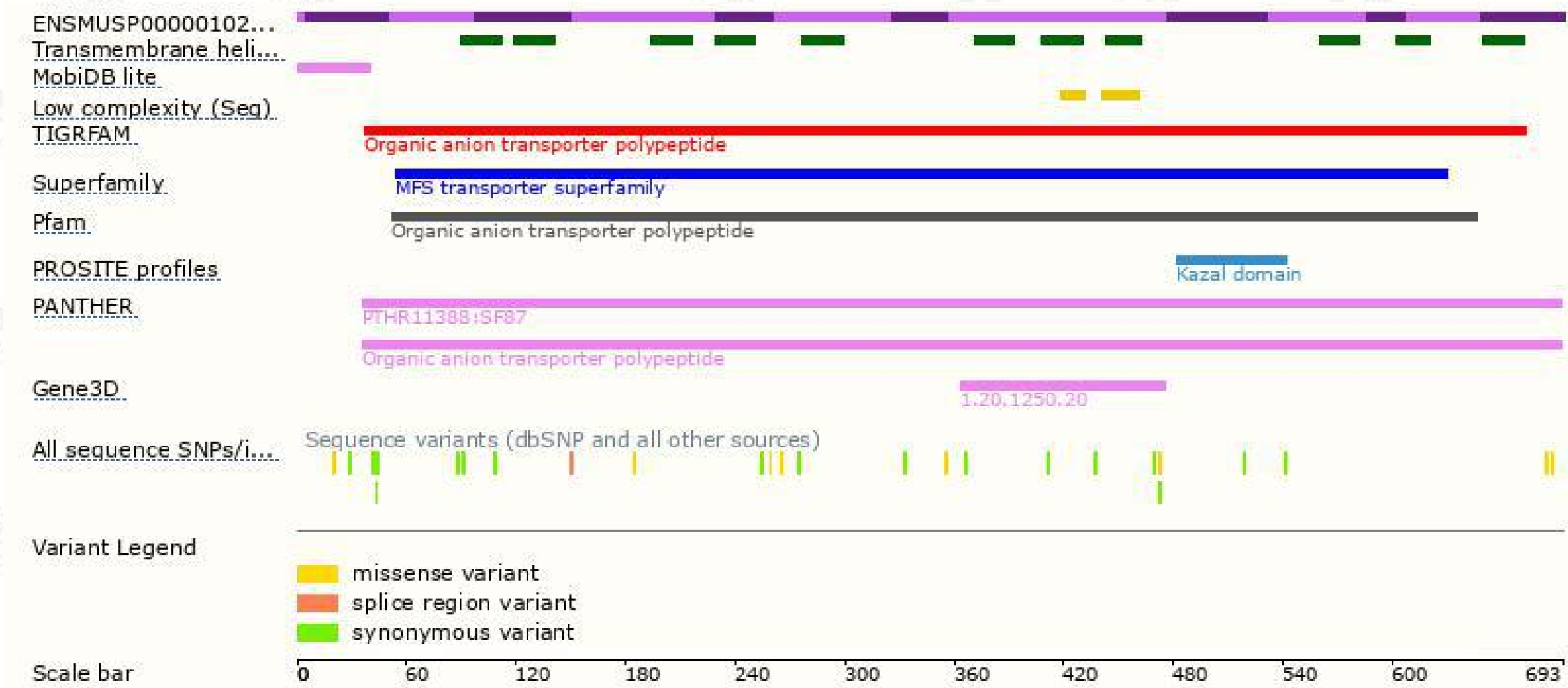
The strategy is based on the design of *Slco2b1-203* transcript,the transcription is shown below:



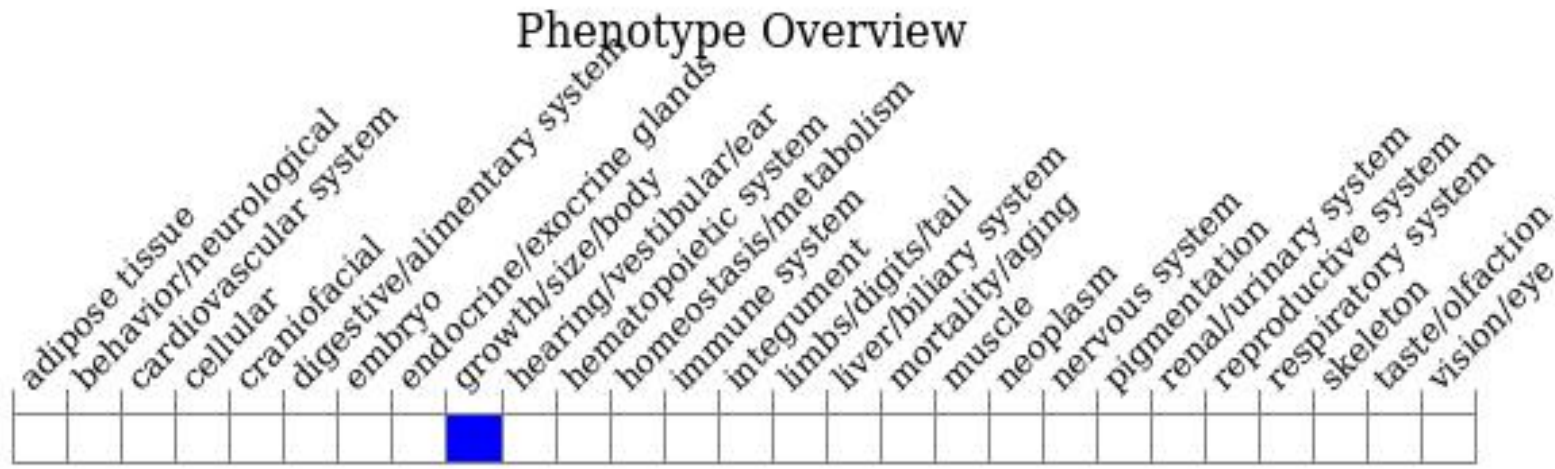
# Genomic location distribution



# 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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