Foxo6 Cas9-KO Strategy

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Design Date: 2019-9-30

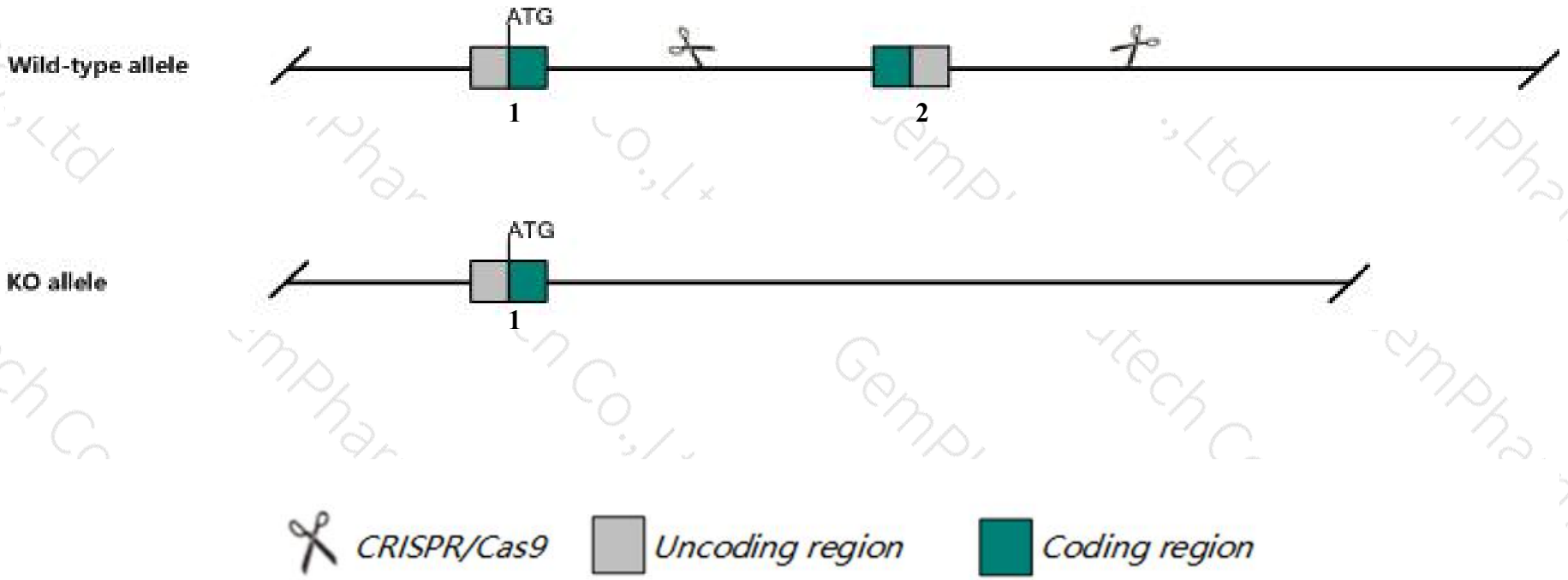
GemPharmatech Co.,Ltd

Project Overview

Project Name	***Foxo6***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Foxo6* gene. The schematic diagram is as follows:



Technical routes

➢ The *Foxo6* gene has 1 transcript. According to the structure of *Foxo6* gene, exon2 of *Foxo6-201* (ENSMUST00000102656.3) transcript is recommended as the knockout region.The region contains most 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 *Foxo6* gene. The brief process is as follows: CRISPR/Cas9 system

Notice

- According to the existing MGI data,Homozygotes for a null allele show defective memory consolidation with impaired neuronal synchronization and altered dendritic spine morphology. Homozygotes for another null allele show attenuated gluconeogenesis, improved glucose tolerance and increased insulin sensitivity after high fat feeding.

- The *Foxo6* gene is located on the Chr4. 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）

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Foxo6 forkhead box O6 [Mus musculus (house mouse)]

Gene ID: 329934, updated on 31-Jan-2019

Summary

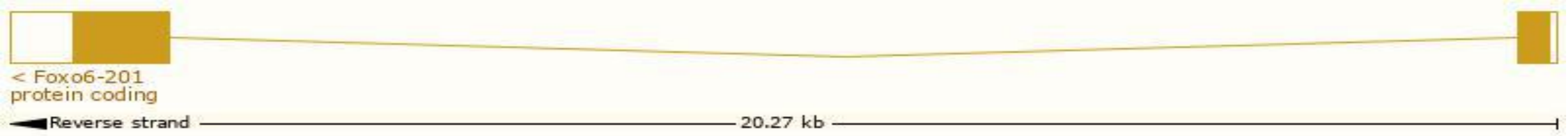
Official Symbol	Foxo6 provided by MGI
Official Full Name	forkhead box O6 provided byMGI
Primary source	MGI:MGI:2676586
See related	Ensembl:ENSMUSG00000052135
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Broad expression in small intestine adult (RPKM 9.5), whole brain E14.5 (RPKM 8.7) and 19 other tissuesSee more
Orthologs	human all

Transcript information（Ensembl）

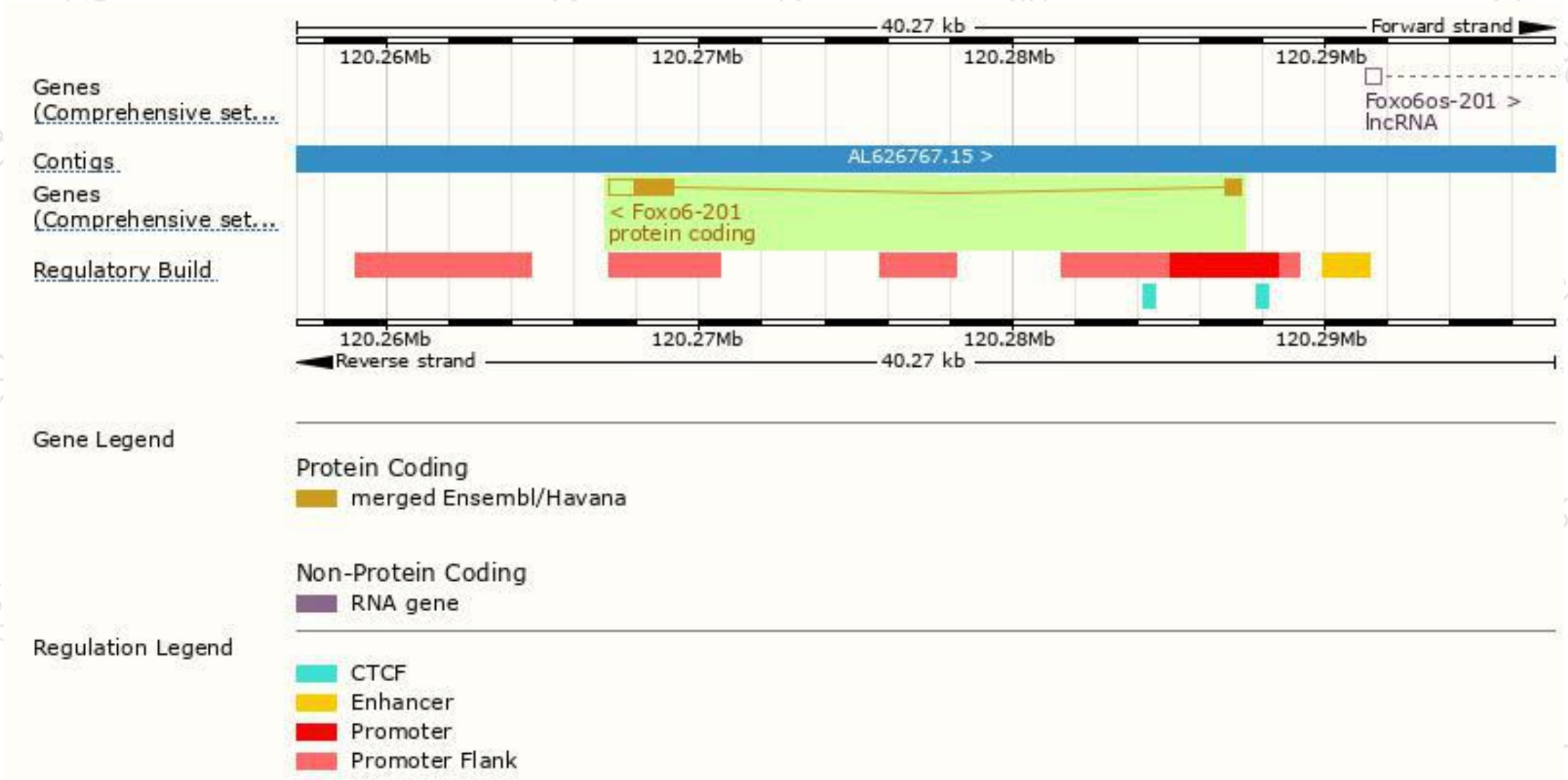
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Foxo6-201	ENSMUST00000102656.3	2615	559aa	Protein coding	CCDS18588	Q70KY4	TSL:1 GENCODE basic APPRIS P1

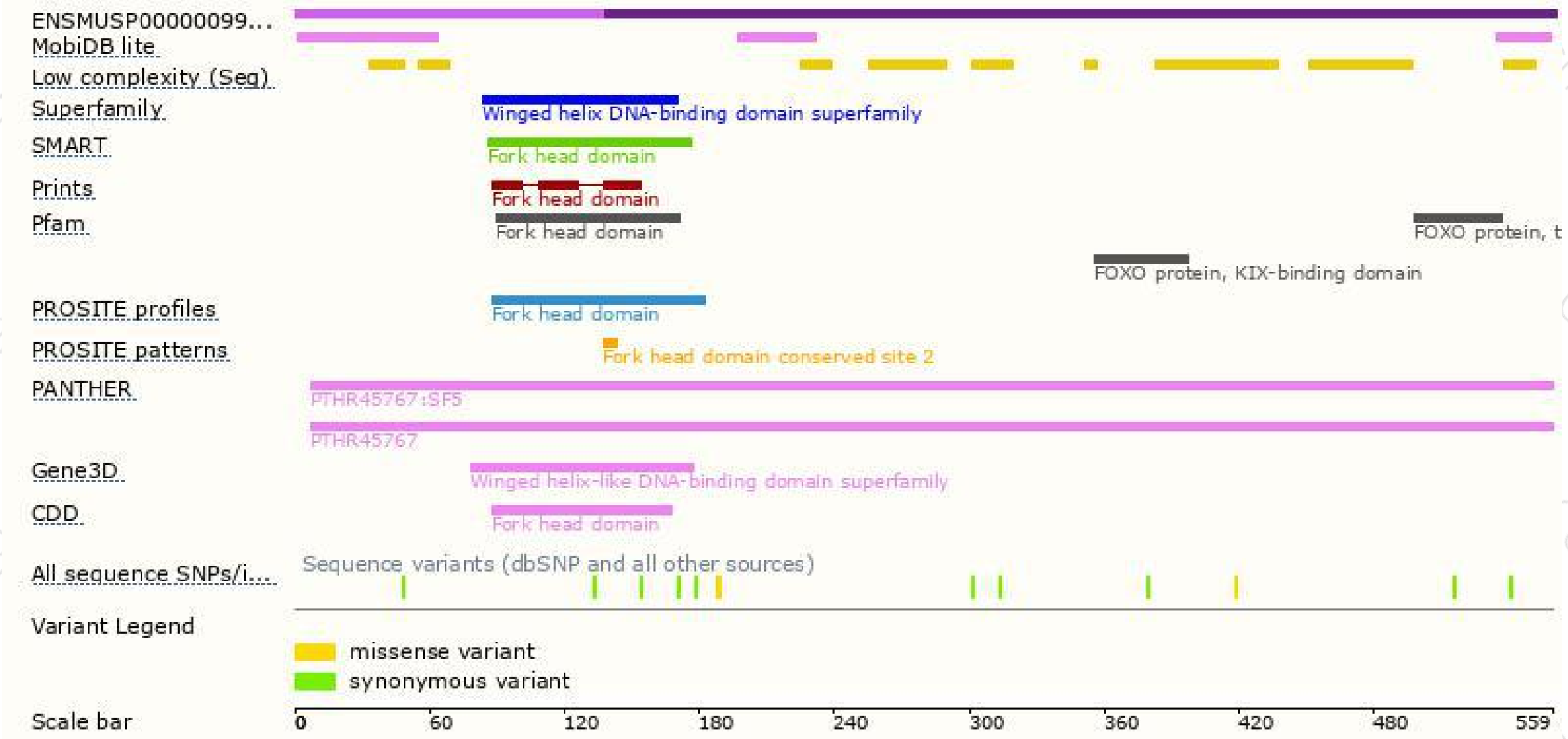
The strategy is based on the design of *Foxo6-201* transcript,The transcription is shown below



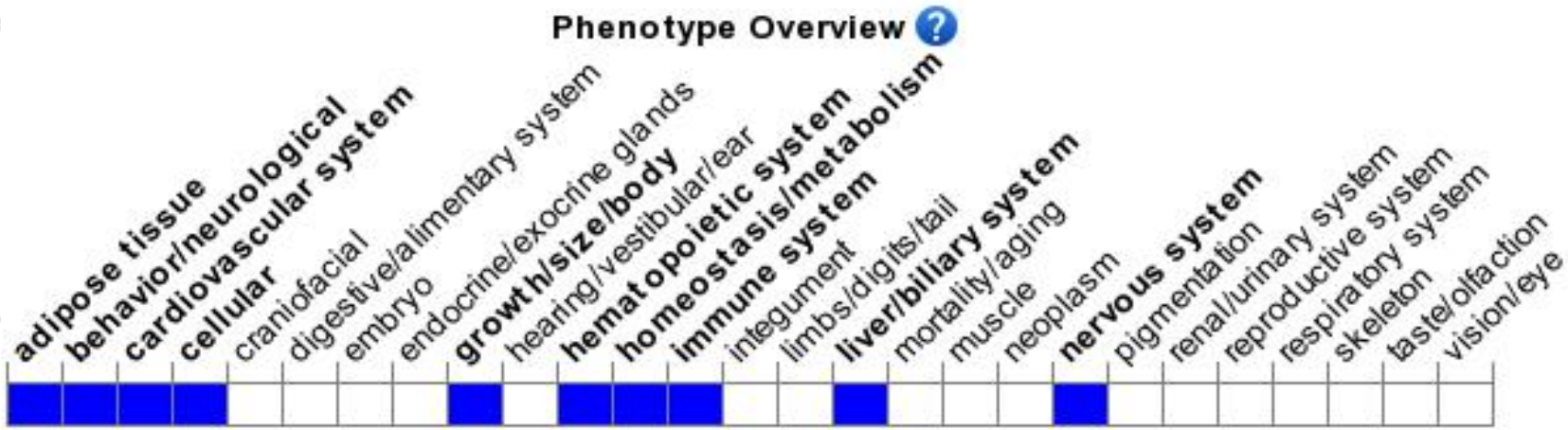
Genomic location distribution



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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