Rab11fip5 Cas9-CKO Strategy

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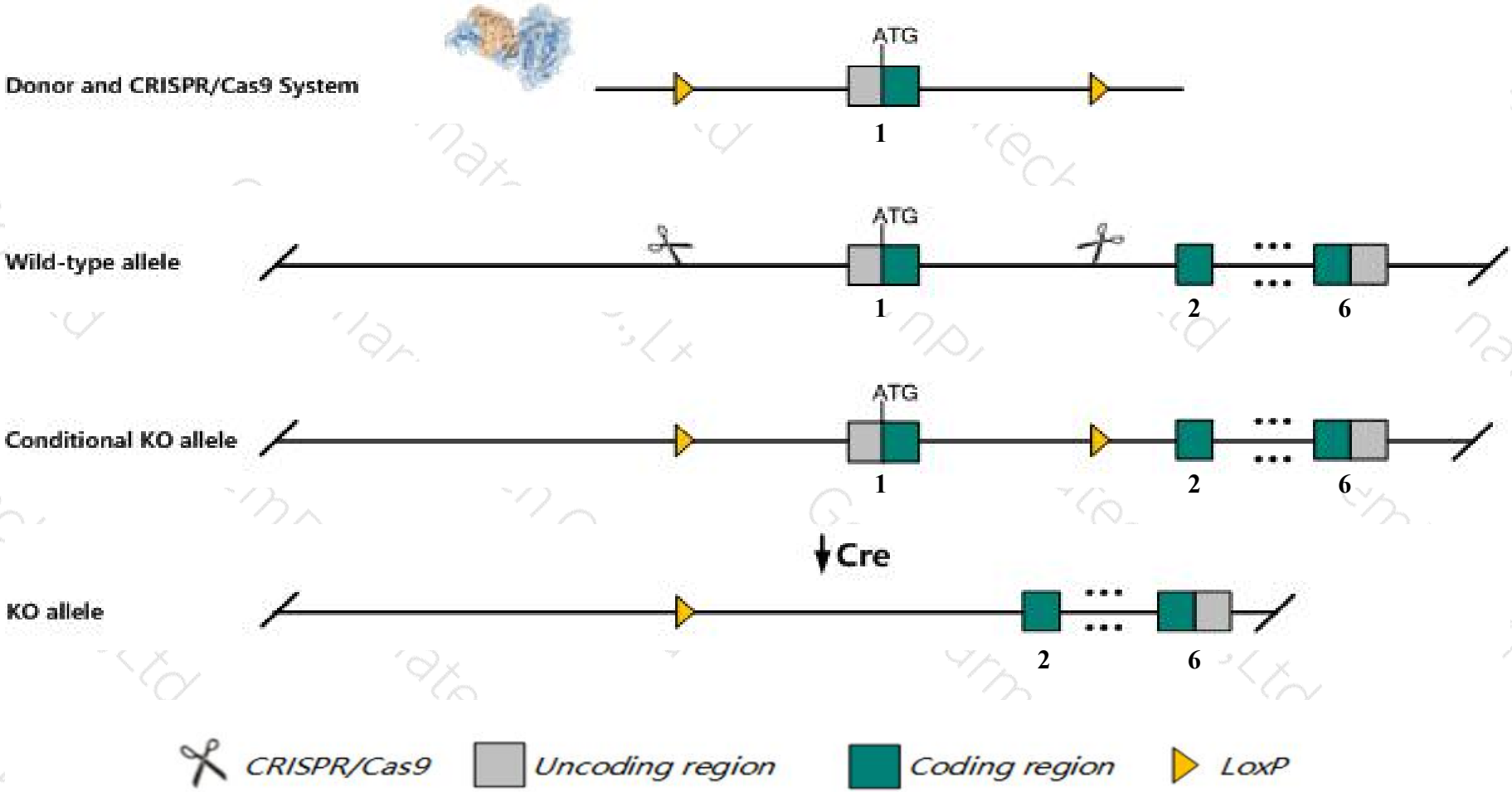
Project Overview

Project Name	***Rab11fip5***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rab11fip5* gene. The schematic diagram is as follows:



Technical routes

- The *Rab11fip5* gene has 3 transcripts. According to the structure of *Rab11fip5* gene, exon1 of *Rab11fip5-202* (ENSMUST00000204087.1) 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 *Rab11fip5* 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 floxed allele are viable and fertile.

➢ The *Rab11fip5* gene is located on the Chr6. 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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Rab11fip5 RAB11 family interacting protein 5 (class I) [Mus musculus (house mouse)]

Gene ID: 52055, updated on 31-Jan-2019

Summary

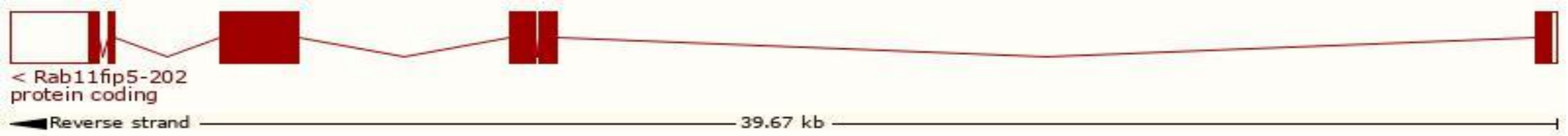
Official Symbol	Rab11fip5 provided by MGI
Official Full Name	RAB11 family interacting protein 5 (class I) provided by MGI
Primary source	MGI:MGI:1098586
See related	Ensembl:ENSMUSG00000051343
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	9130206P09Rik, C75969, D6Ertd32e, GAF1, RIP11
Expression	Broad expression in testis adult (RPKM 59.8), ovary adult (RPKM 19.7) and 20 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

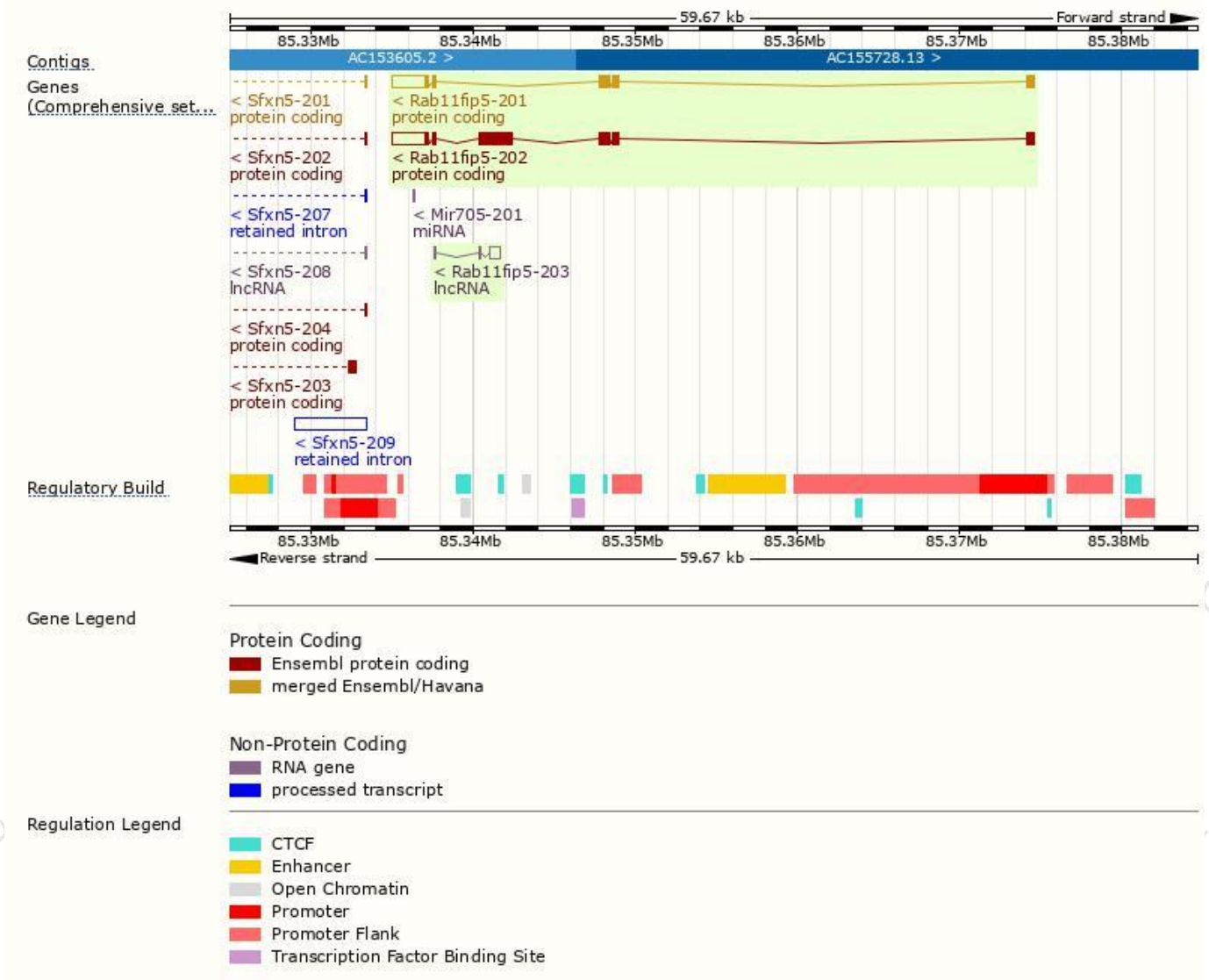
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rab11fip5-202	ENSMUST00000204087.1	6119	1318aa	Protein coding	CCDS85083	A0A0N4SW73	TSL:1 GENCODE basic APPRIS ALT2
Rab11fip5-201	ENSMUST00000060837.9	4035	645aa	Protein coding	CCDS20292	Q8R361	TSL:1 GENCODE basic APPRIS P3
Rab11fip5-203	ENSMUST00000204504.1	784	No protein	lncRNA	-	-	TSL:5

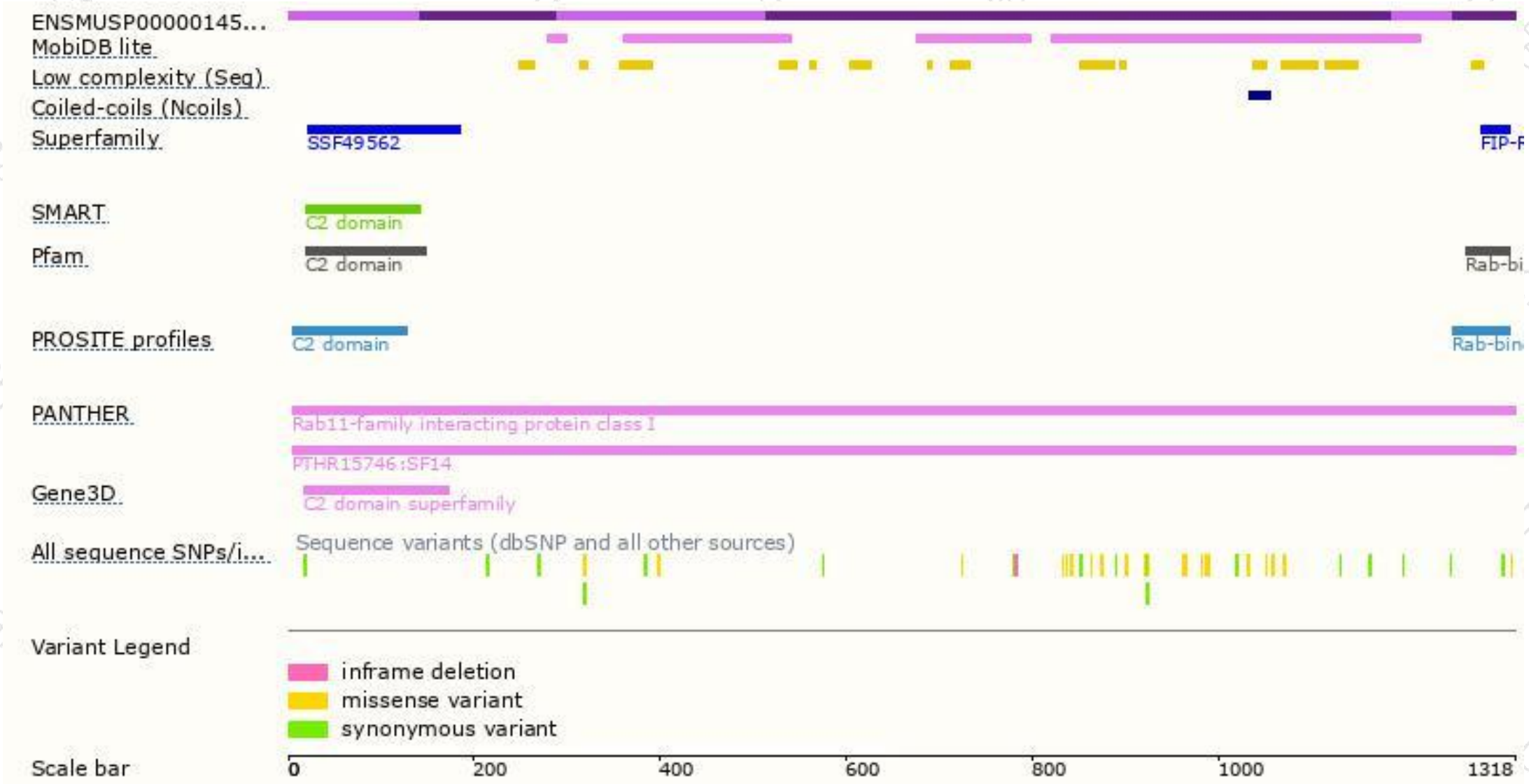
The strategy is based on the design of *Rab11fip5-202* transcript,The transcription is shown below



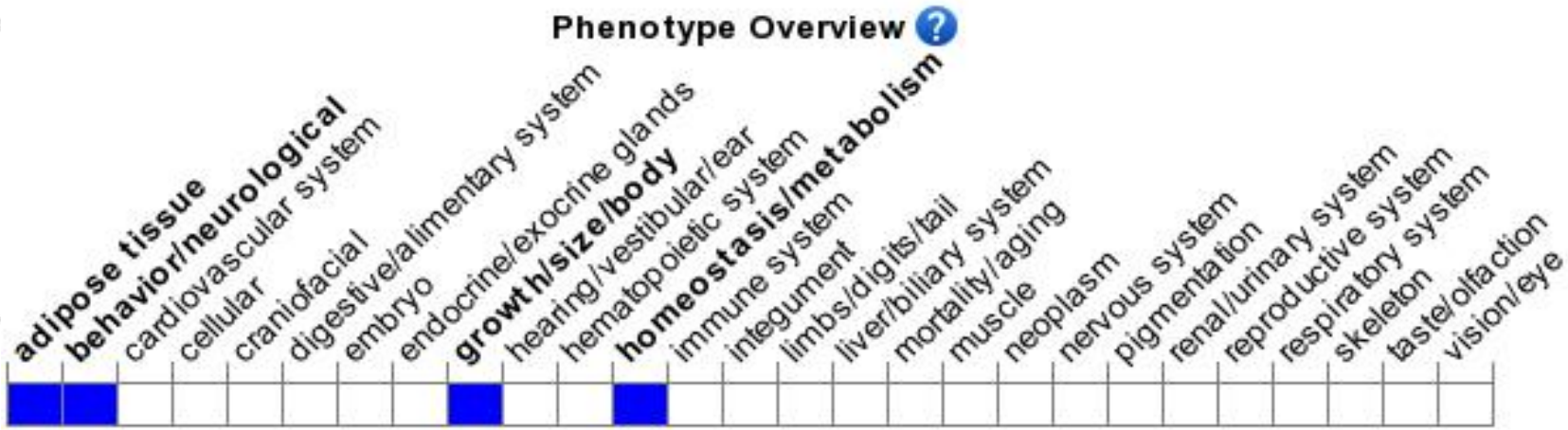
Genomic location distribution



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/).

According to the existing MGI data,Mice homozygous for a floxed allele are viable and fertile.

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If you have any questions, you are welcome to inquire.

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