Npffr1 Cas9-CKO Strategy

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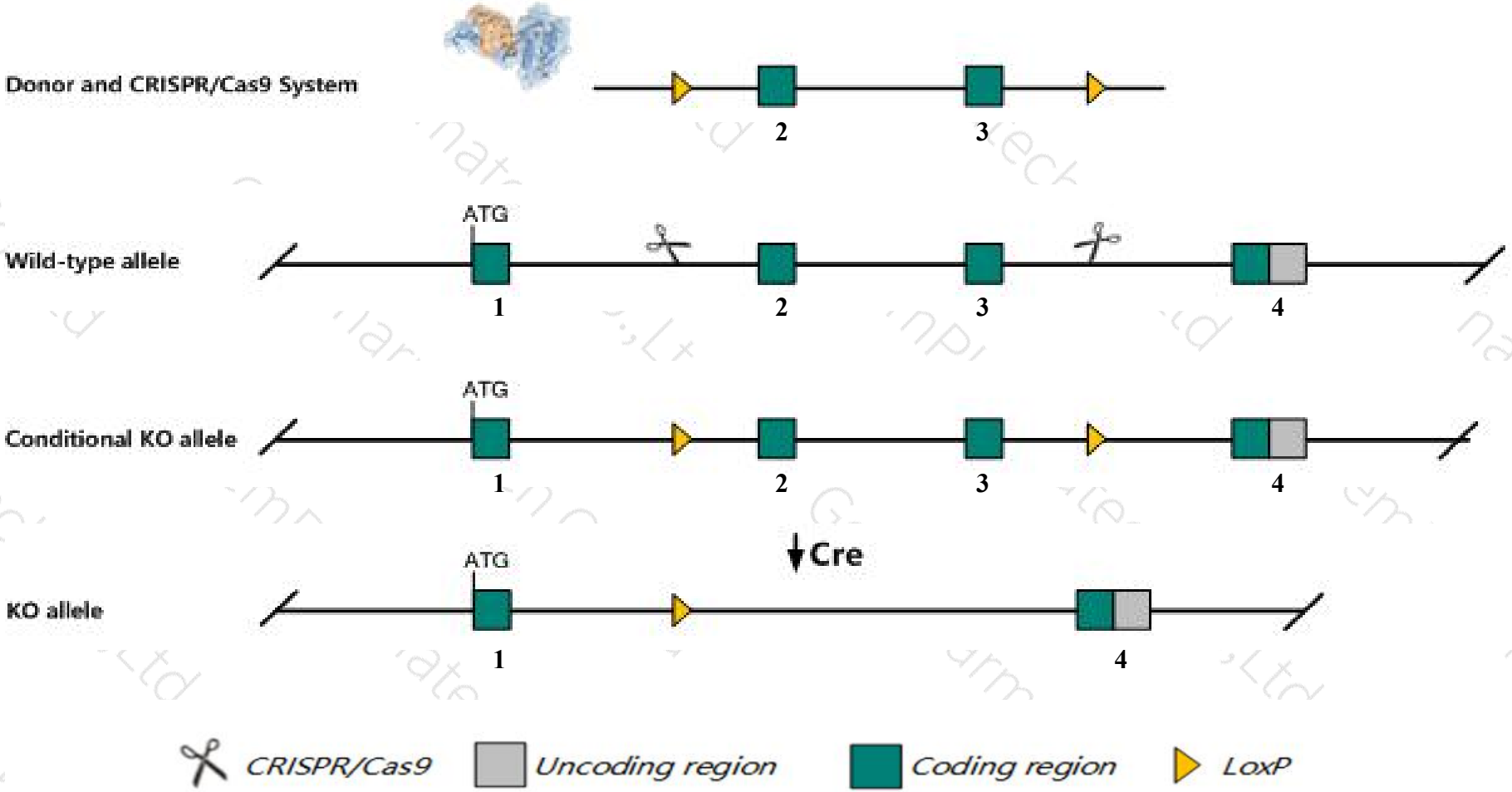
Project Overview

Project Name	***Npffr1***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Npffr1* gene. The schematic diagram is as follows:



Technical routes

- The *Npffr1* gene has 1 transcript. According to the structure of *Npffr1* gene, exon2-exon3 of *Npffr1-201* (ENSMUST00000020287.7) transcript is recommended as the knockout region.The region contains 415bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Npffr1* 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 mutation display abnormal pituitary function with abnormal levels of follicle stimulating and luteinizing hormone levels and increased litter sizes.
- The *Npffr1* gene is located on the Chr10. 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.
- The CKO region contains functional region of the Gm28447 gene.Knockout the region may affect the function of Gm28447 gene.
- 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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Npffr1 neuropeptide FF receptor 1 [Mus musculus (house mouse)]

Gene ID: 237362, updated on 2-Apr-2019

Summary

Official Symbol	Npffr1 provided by MGI
Official Full Name	neuropeptide FF receptor 1 provided byMGI
Primary source	MGI:MGI:2685082
See related	Ensembl:ENSMUSG00000020090
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	Gm236, Gpr147, NPFF1, NPFF1R, OT7T022
Expression	Low expression observed in reference datasetSee more
Orthologs	human all

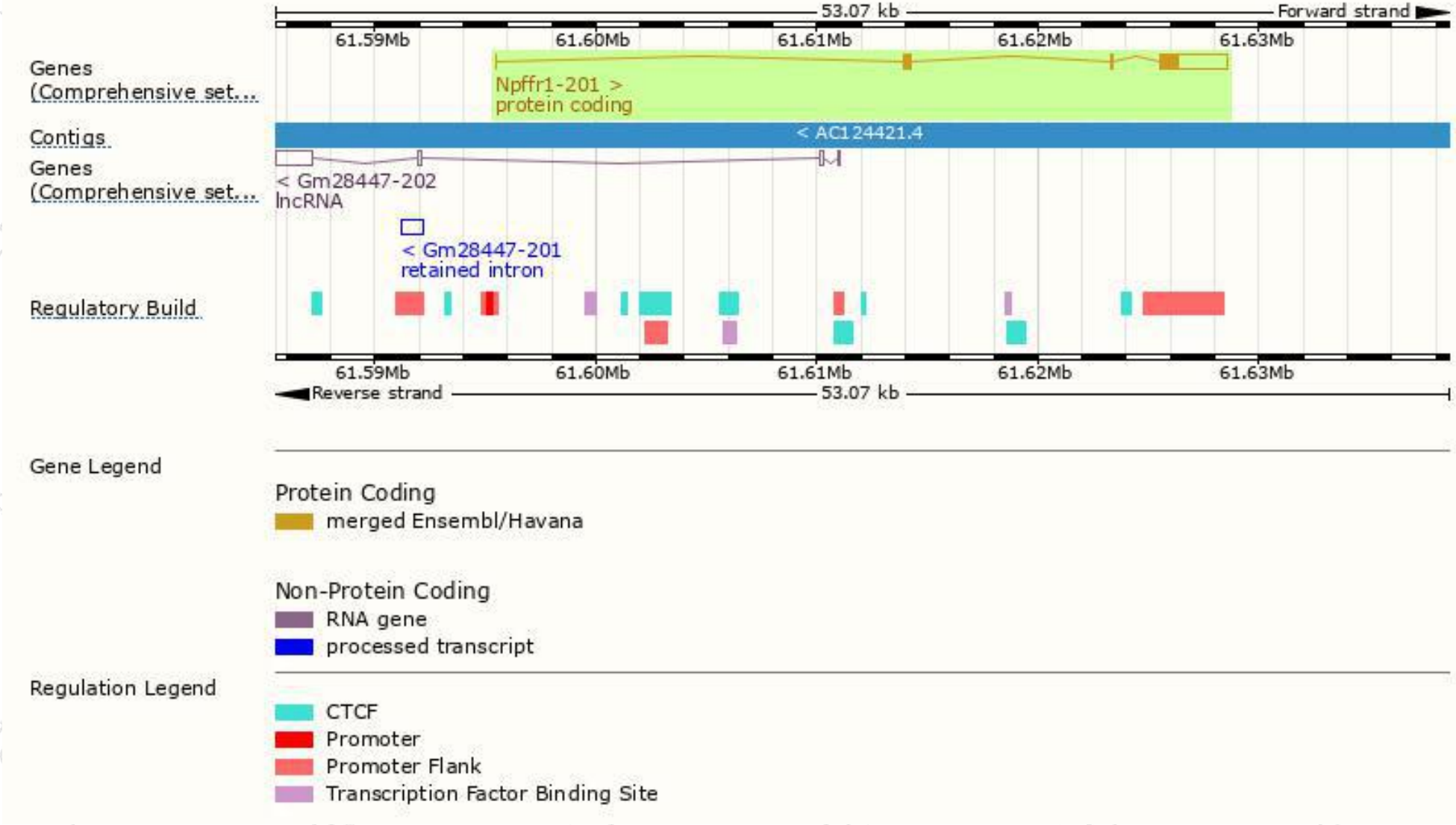
Transcript information（Ensembl）

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

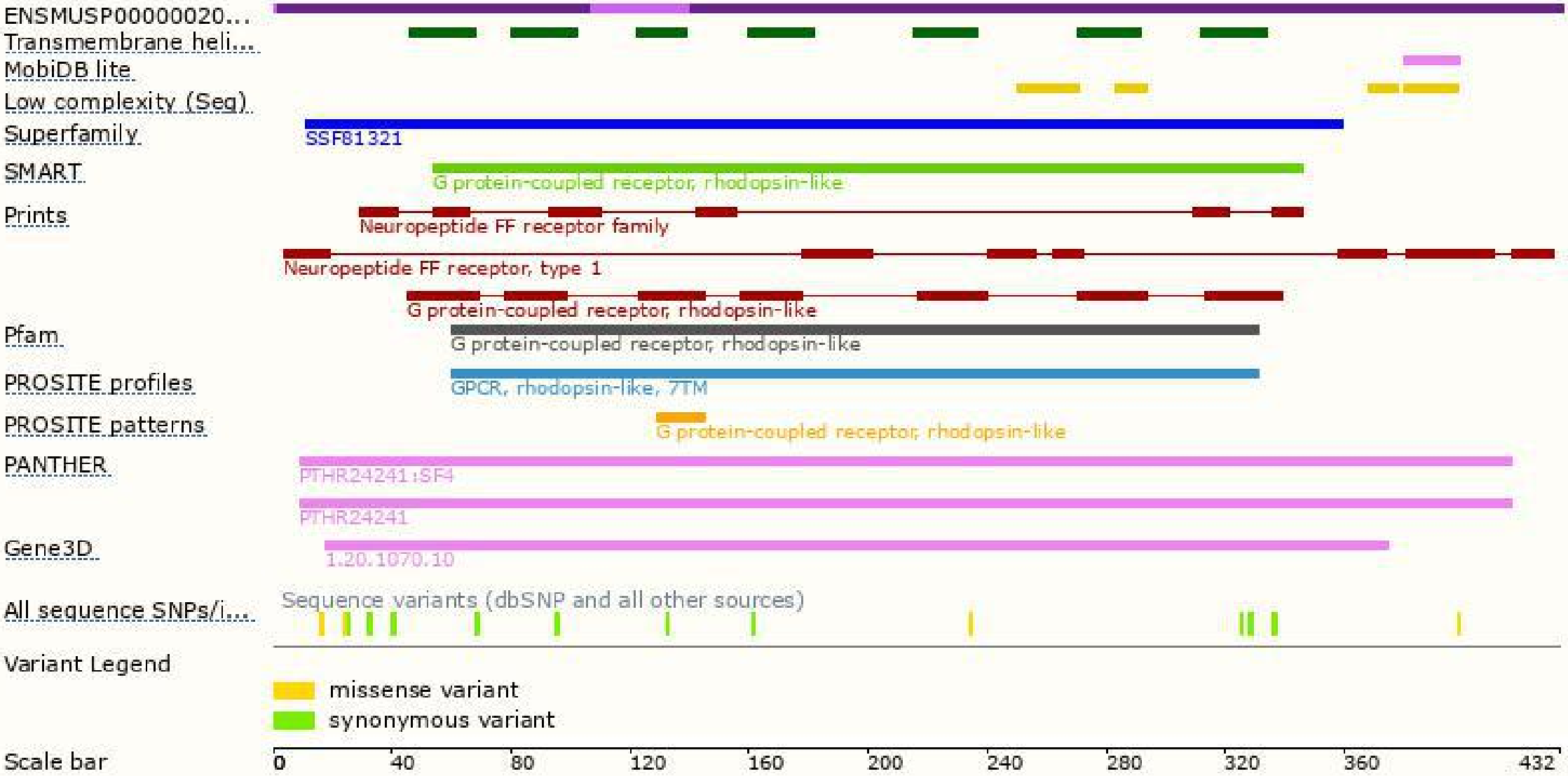
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Npffr1-201	ENSMUST00000020287.7	3479	432aa	Protein coding	CCDS48574	E9Q468	TSL:5 GENCODE basic APPRIS P1

The strategy is based on the design of *Npffr1-201* transcript,The transcription is shown below

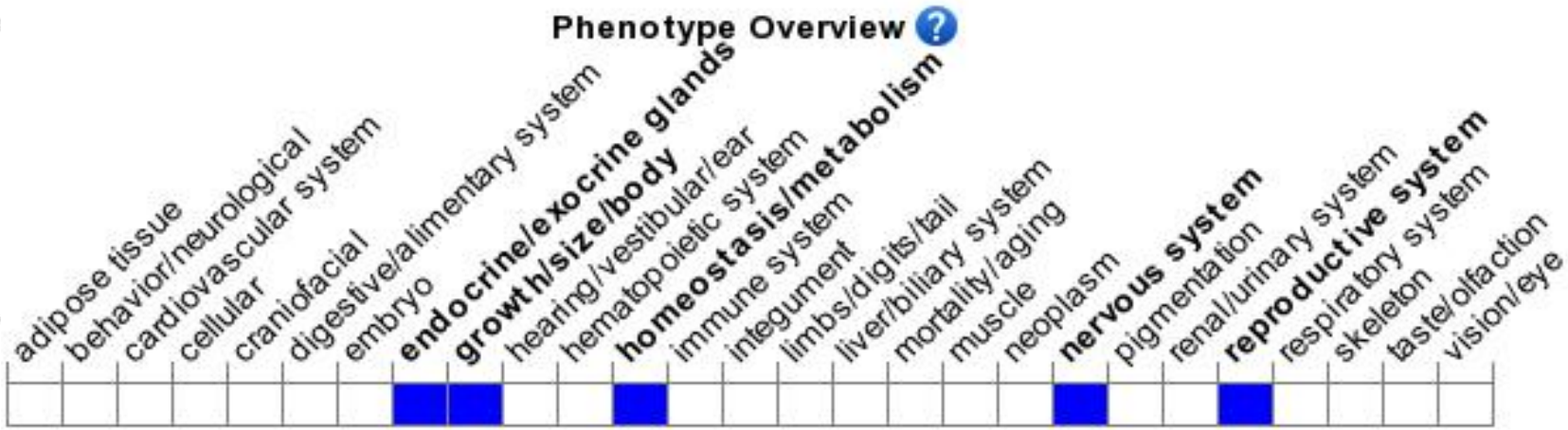
Genomic location distribution



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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