Rc3h1 Cas9-CKO Strategy

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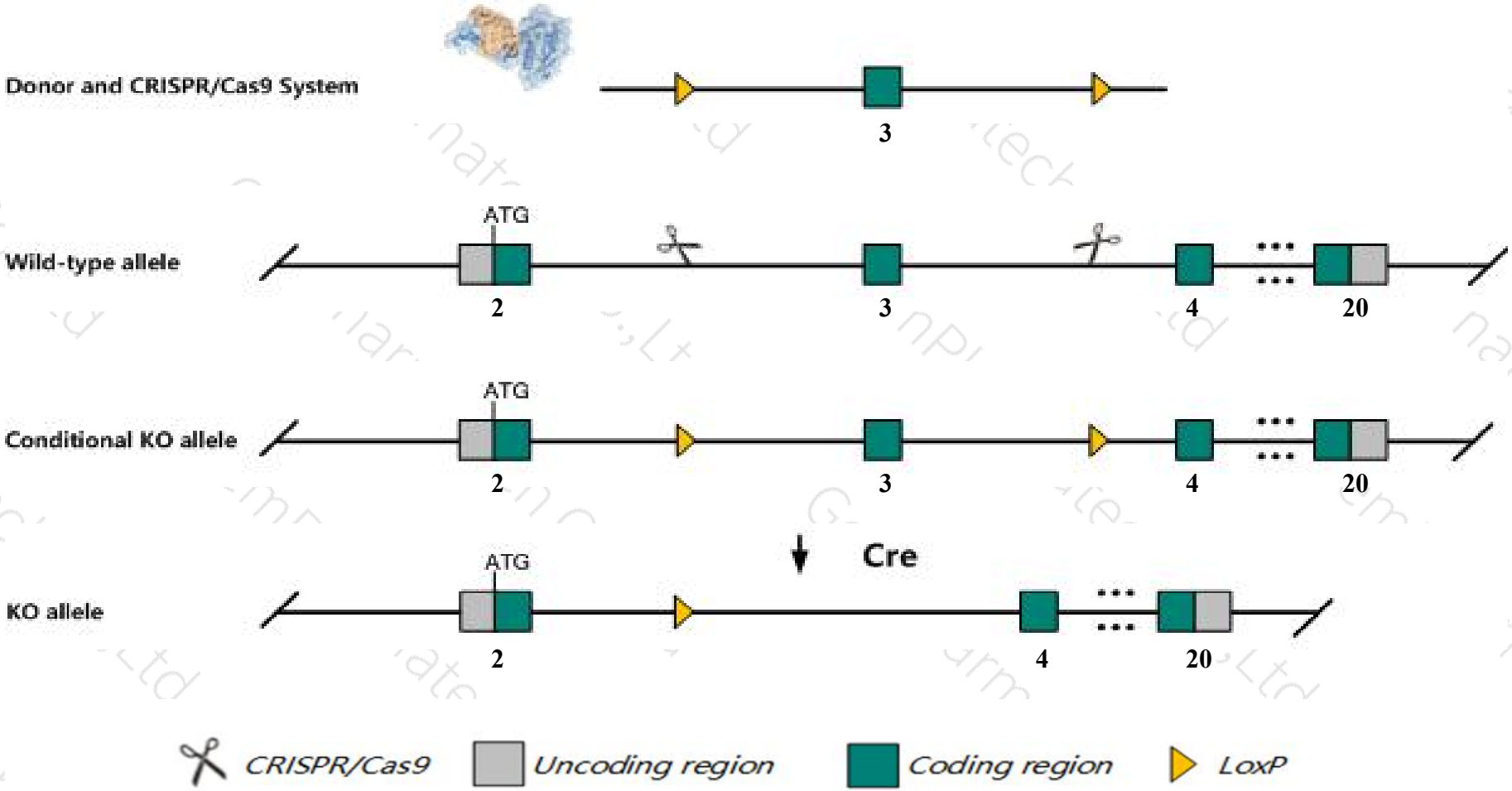
Project Overview

Project Name	***Rc3h1***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rc3h1* gene. The schematic diagram is as follows:



Technical routes

- The *Rc3h1* gene has 3 transcripts. According to the structure of *Rc3h1* gene, exon3 of *Rc3h1-202* (ENSMUST00000161609.7) transcript is recommended as the knockout region.The region contains 121bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Rc3h1* 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,A single recessive mutation on this gene resulted in severe autoimmune disease with phenotype resembling human systemic lupus erythematosus.

- The *Rc3h1* gene is located on the Chr1. 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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Rc3h1 RING CCCH (C3H) domains 1 [Mus musculus (house mouse)]

Gene ID: 381305, updated on 31-Jan-2019

Summary

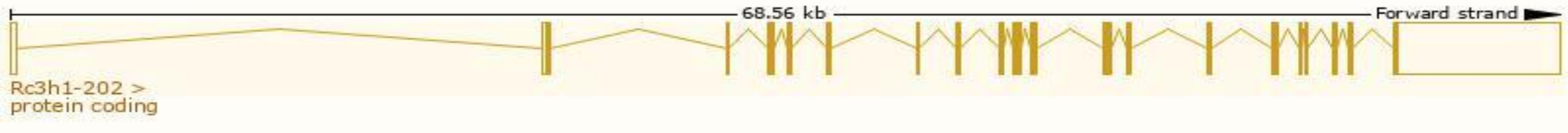
Official Symbol	Rc3h1 provided by MGI
Official Full Name	RING CCCH (C3H) domains 1 provided byMGI
Primary source	MGI:MGI:2685397
See related	Ensembl:ENSMUSG00000040423
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	5730557L09Rik, Gm551, N28103, mKIAA2025
Expression	Ubiquitous expression in testis adult (RPKM 5.7), thymus adult (RPKM 5.0) and 28 other tissuesSee more
Orthologs	human all

Transcript information（Ensembl）

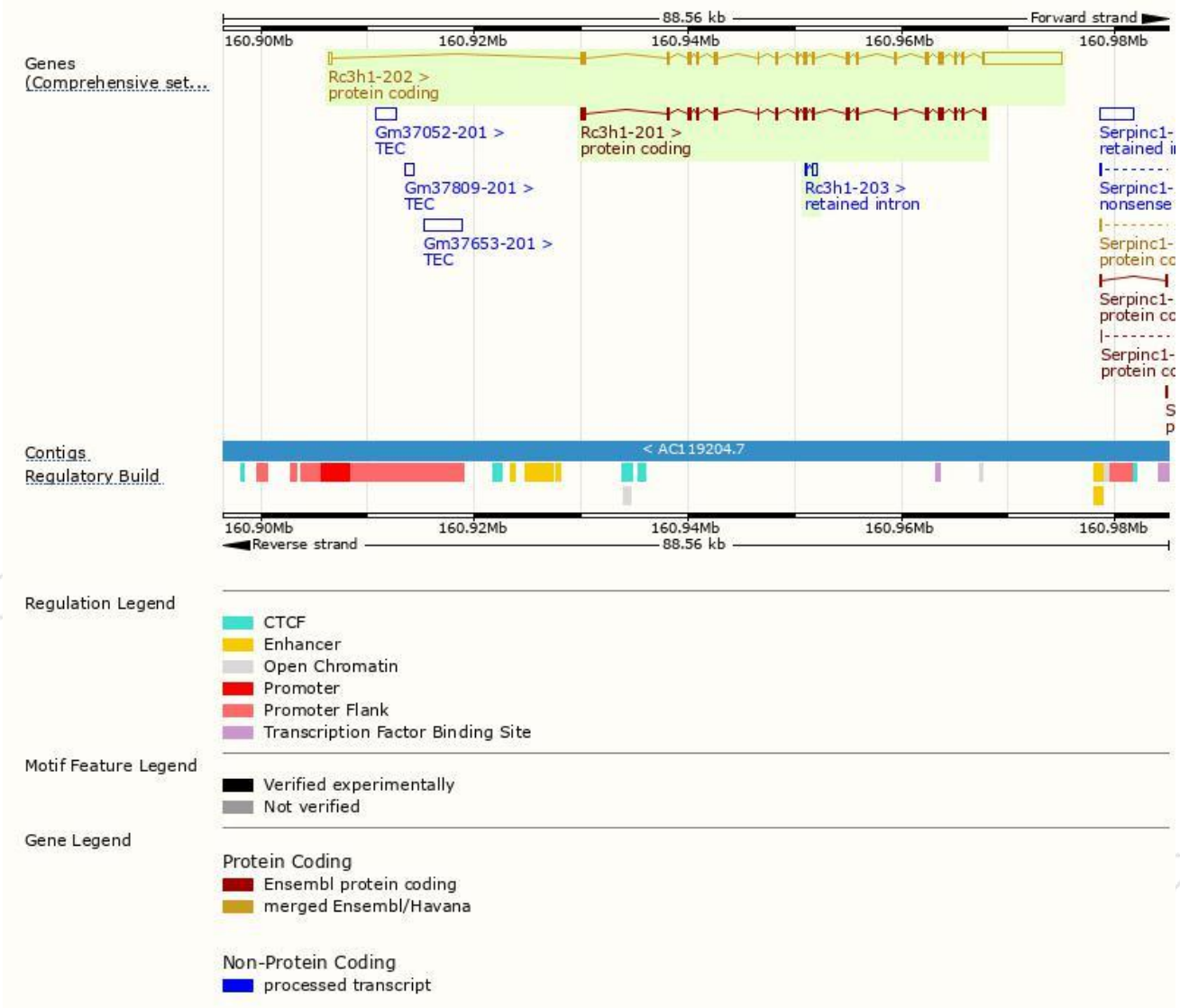
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rc3h1-202	ENSMUST00000161609.7	11006	1130aa	Protein coding	CCDS15410	Q4VGL6	TSL:1 GENCODE basic APPRIS P2
Rc3h1-201	ENSMUST00000035911.4	3476	1121aa	Protein coding	-	H7BX02	TSL:5 GENCODE basic APPRIS ALT2
Rc3h1-203	ENSMUST00000161708.1	663	No protein	Retained intron	-	-	TSL:3

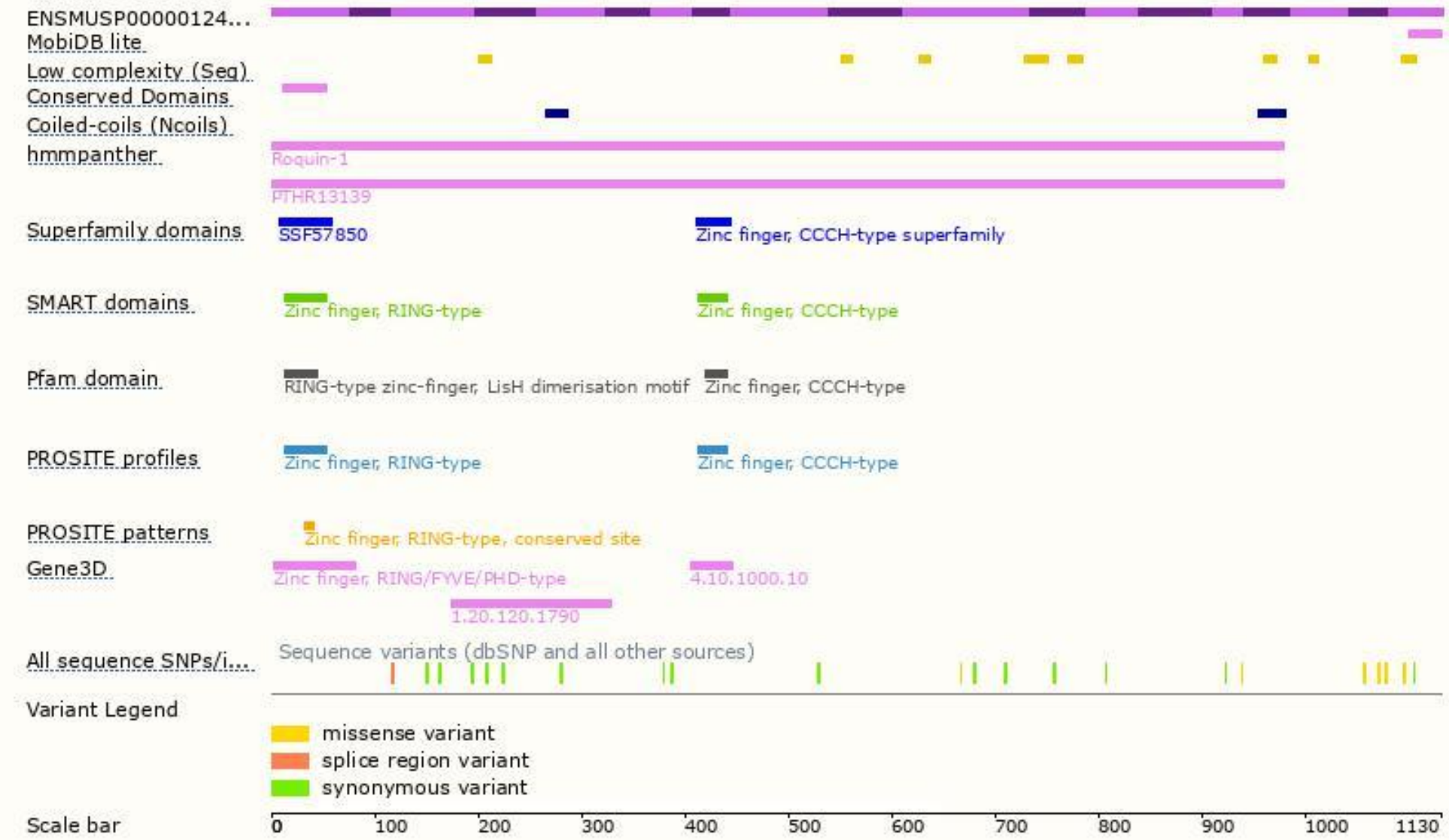
The strategy is based on the design of *Rc3h1-202* transcript,The transcription is shown below



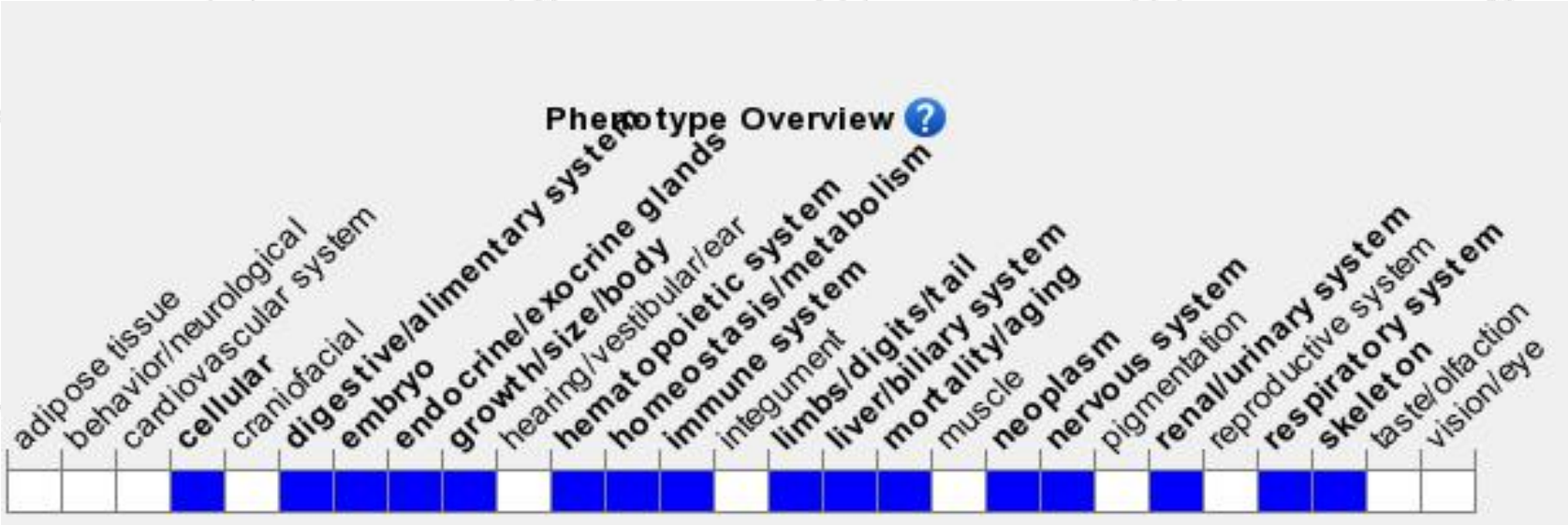
Genomic location distribution



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/).

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

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