Kdr Cas9-CKO Strategy

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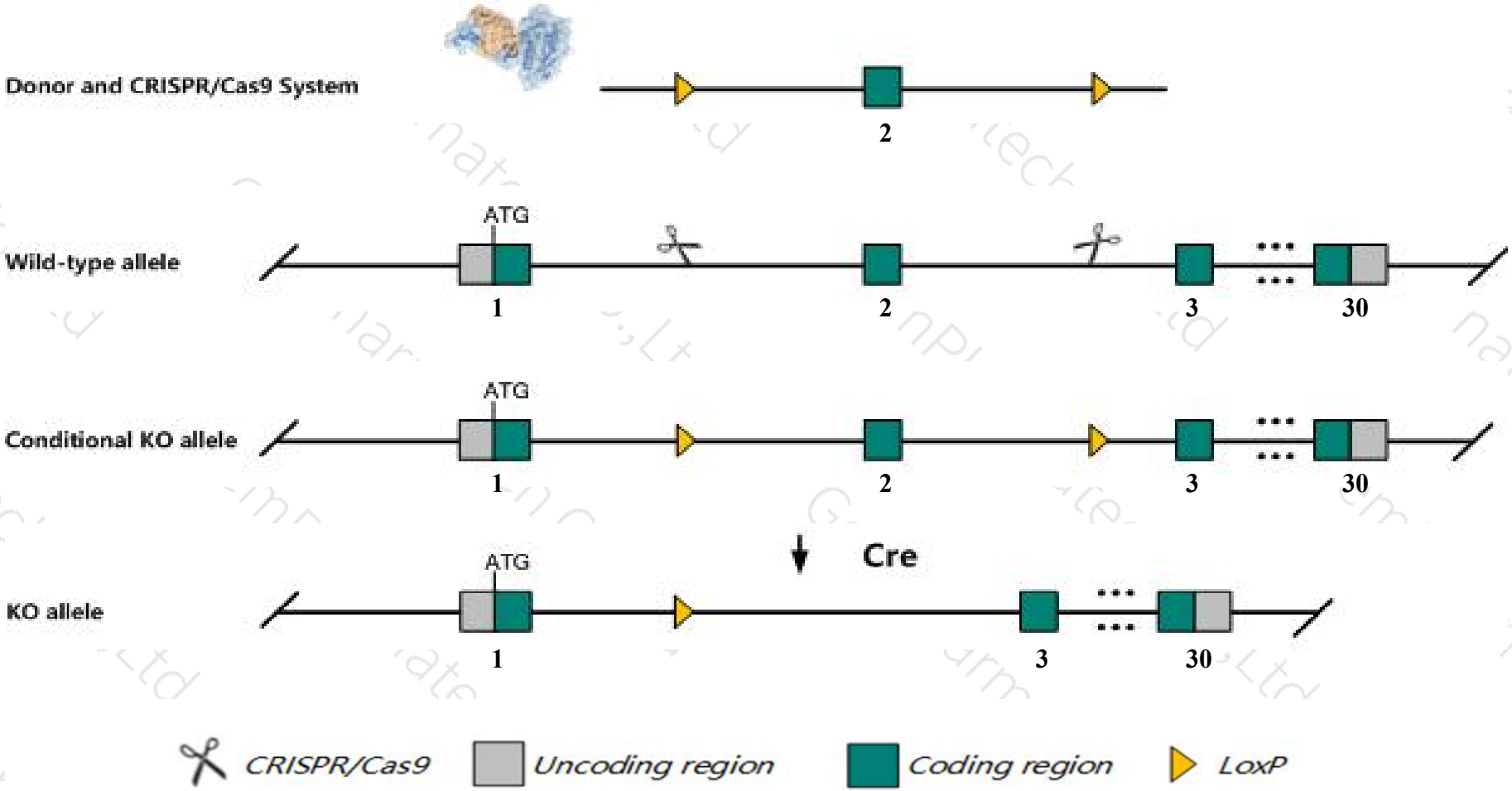
Project Overview

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

Conditional Knockout strategy

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



Technical routes

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- The *Kdr* gene has 3 transcripts. According to the structure of *Kdr* gene, exon2 of *Kdr-201* (ENSMUST00000113516.1) transcript is recommended as the knockout region.The region contains 94bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Kdr* 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,Homozygous mice die at early embryonic stages due to failure of blood vessel formation.

- The *Kdr* gene is located on the Chr5. 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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Kdr kinase insert domain protein receptor [Mus musculus (house mouse)]

Gene ID: 16542, updated on 9-Apr-2019

Summary

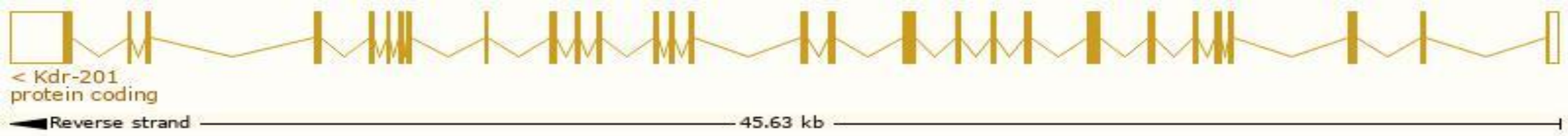
Official Symbol	Kdr provided by MGI
Official Full Name	kinase insert domain protein receptor provided by MGI
Primary source	MGI:MGI:96683
See related	Ensembl:ENSMUSG00000062960
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	6130401C07, Flk-1, Flk1, Krd-1, Ly73, VEGFR-2, VEGFR2, orv, sVEGFR-2
Expression	Broad expression in lung adult (RPKM 42.4), heart adult (RPKM 19.3) and 22 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

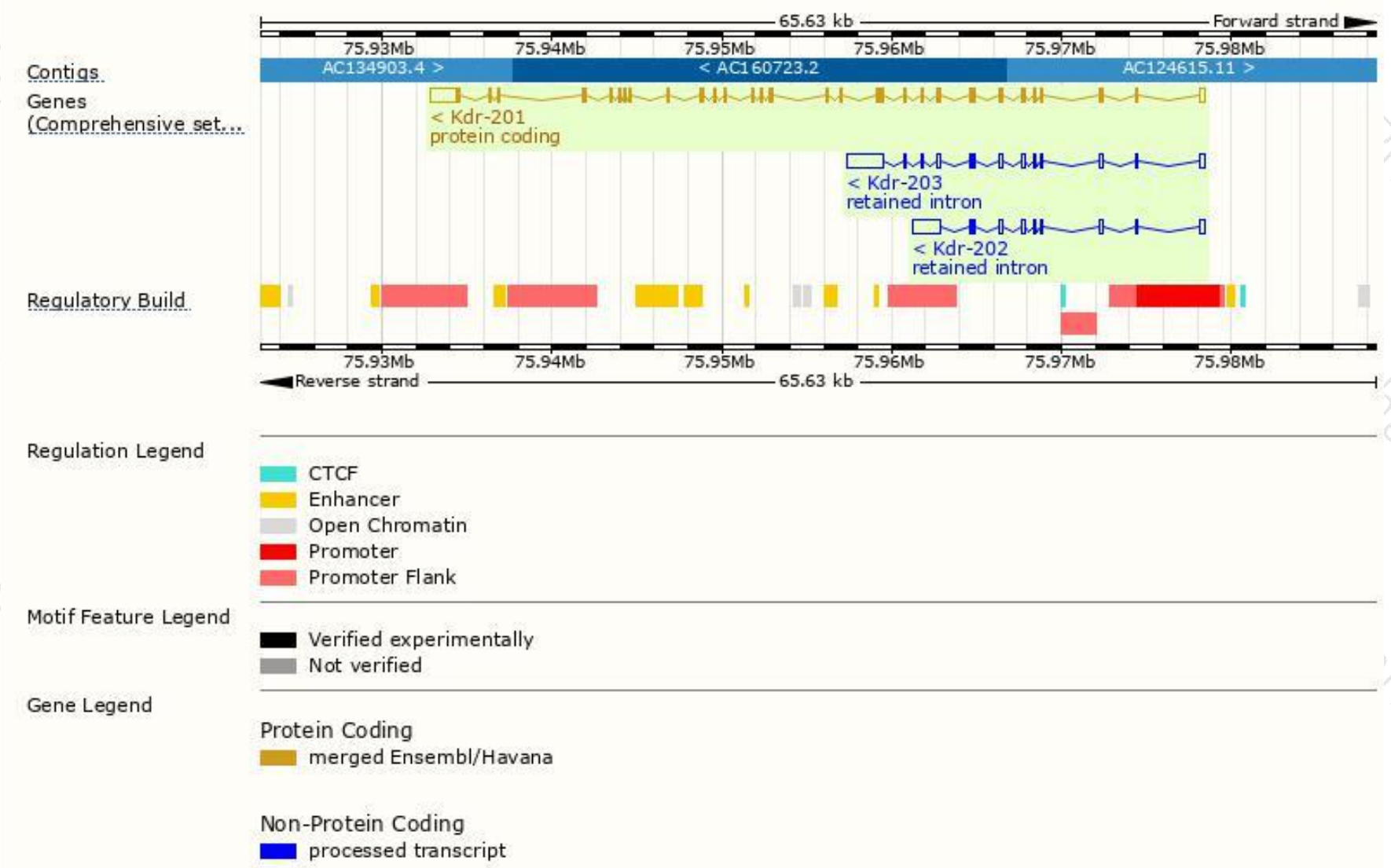
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kdr-201	ENSMUST00000113516.1	5924	1345aa	Protein coding	CCDS39114	Q8VCD0	TSL:1 GENCODE basic APPRIS P1
Kdr-203	ENSMUST00000202473.3	4023	No protein	Retained intron	-	-	TSL:1
Kdr-202	ENSMUST00000149573.1	3174	No protein	Retained intron	-	-	TSL:1

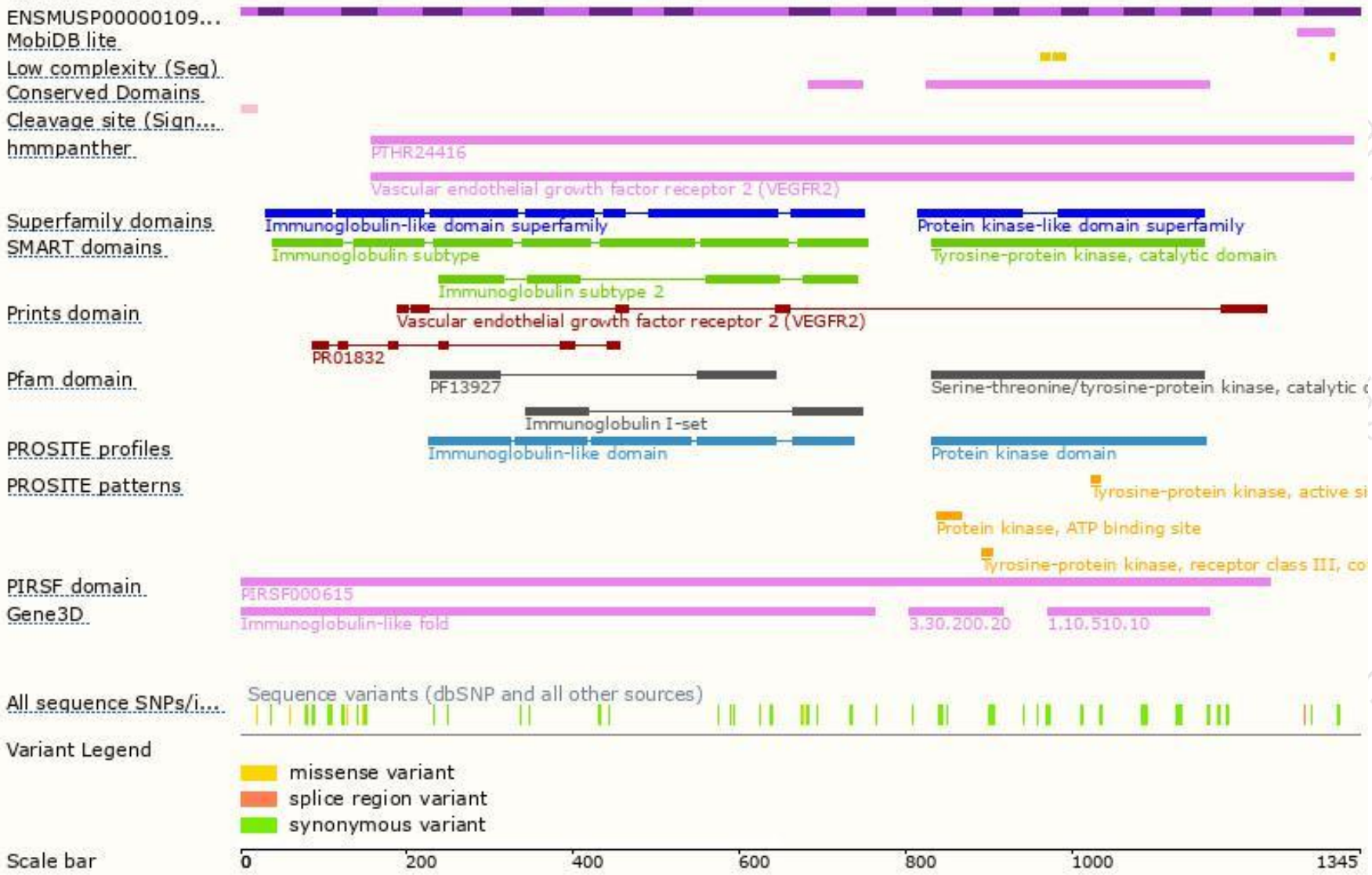
The strategy is based on the design of *Kdr-201* transcript,The transcription is shown below



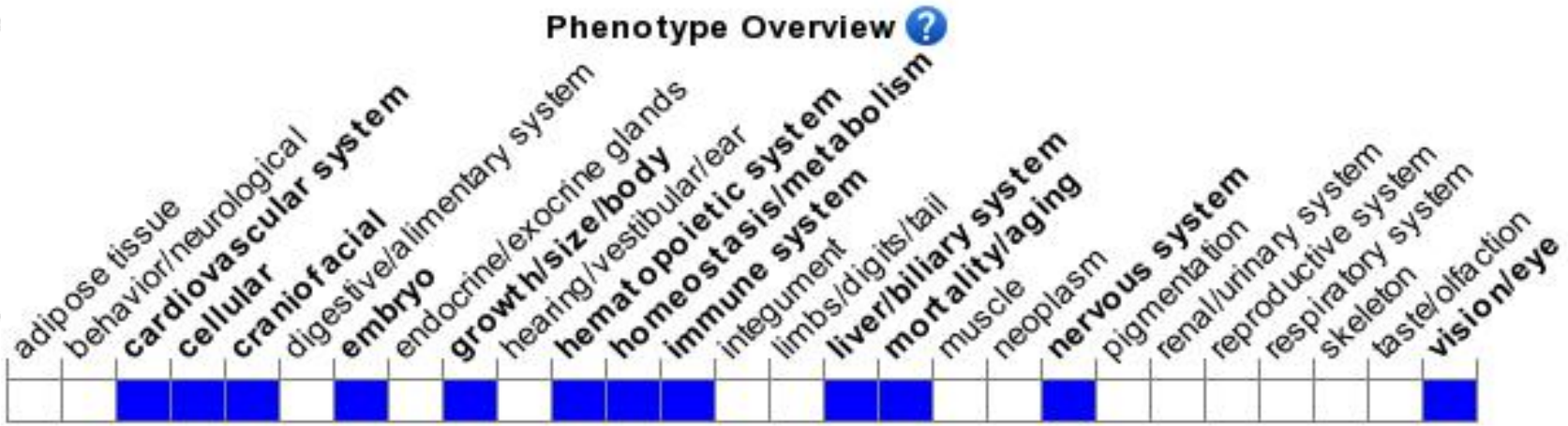
Genomic location distribution



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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