

Vwa1 Cas9-CKO Strategy

Designer:

JiaYu

Reviewer:

Xiaojing Li

Design Date:

2020-3-3

Project Overview

Project Name

Vwa1

Project type

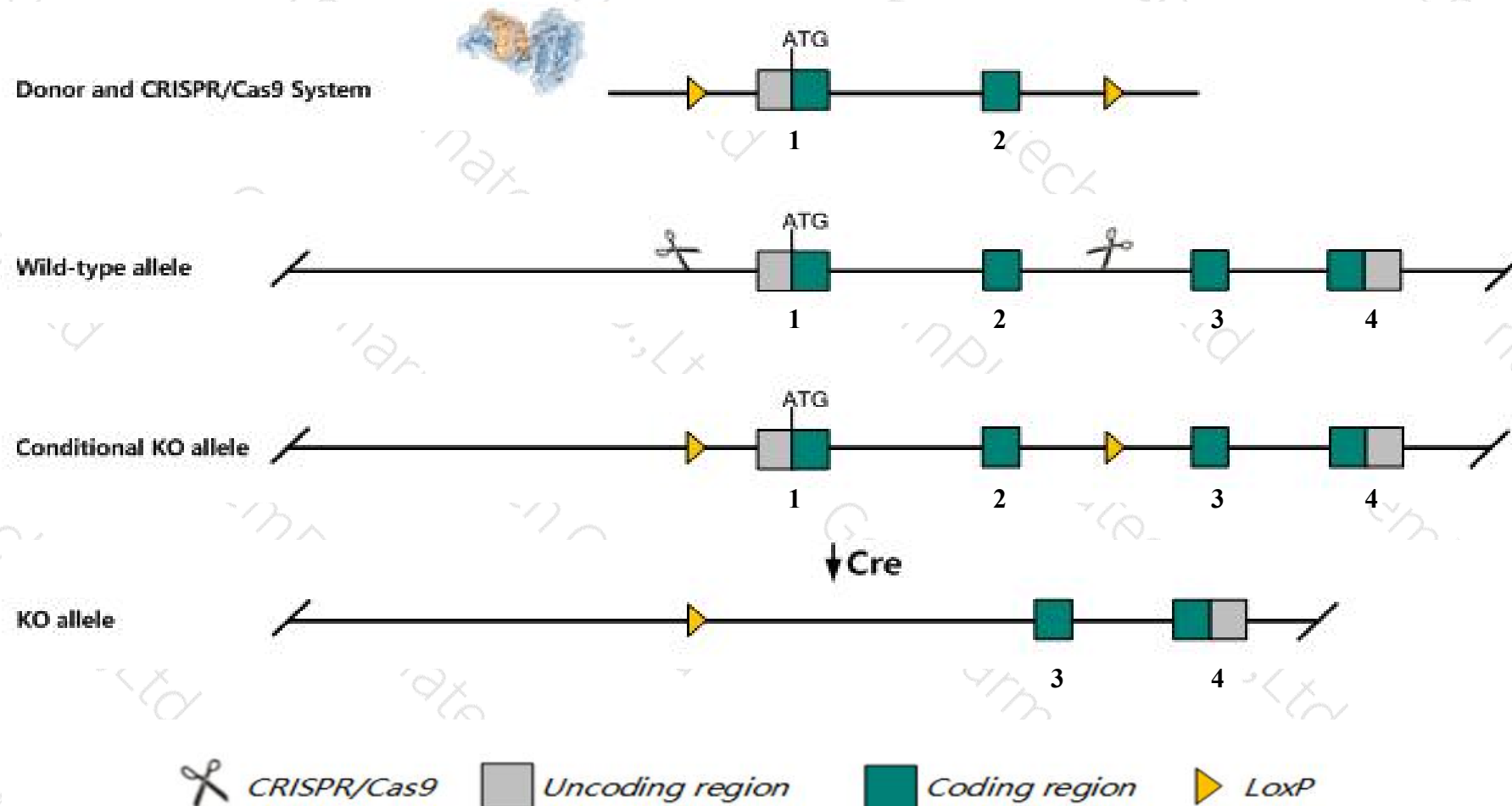
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Vwa1* gene. The schematic diagram is as follows:



Technical routes

- The *Vwal* gene has 1 transcript. According to the structure of *Vwal* gene, exon1-exon2 of *Vwal*-201 (ENSMUST00000042196.3) 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 *Vwal* 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.

- According to the existing MGI data, Mice homozygous for a reporter allele exhibit abnormal motor coordination/balance, increased thermal nociceptive threshold, and altered peripheral nerve structure and function.
- The *Vwal* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Vwa1 von Willebrand factor A domain containing 1 [Mus musculus (house mouse)]

Gene ID: 246228, updated on 3-Feb-2019

Summary



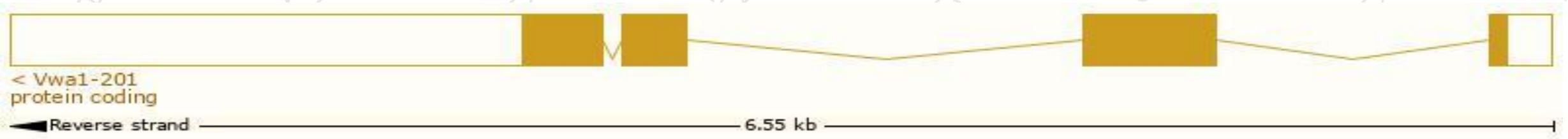
Official Symbol	Vwa1 provided by MGI
Official Full Name	von Willebrand factor A domain containing 1 provided by MGI
Primary source	MGI:MGI:2179729
See related	Ensembl:ENSMUSG00000042116
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4932416A11Rik, WARP
Expression	Broad expression in duodenum adult (RPKM 40.7), small intestine adult (RPKM 34.4) and 23 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

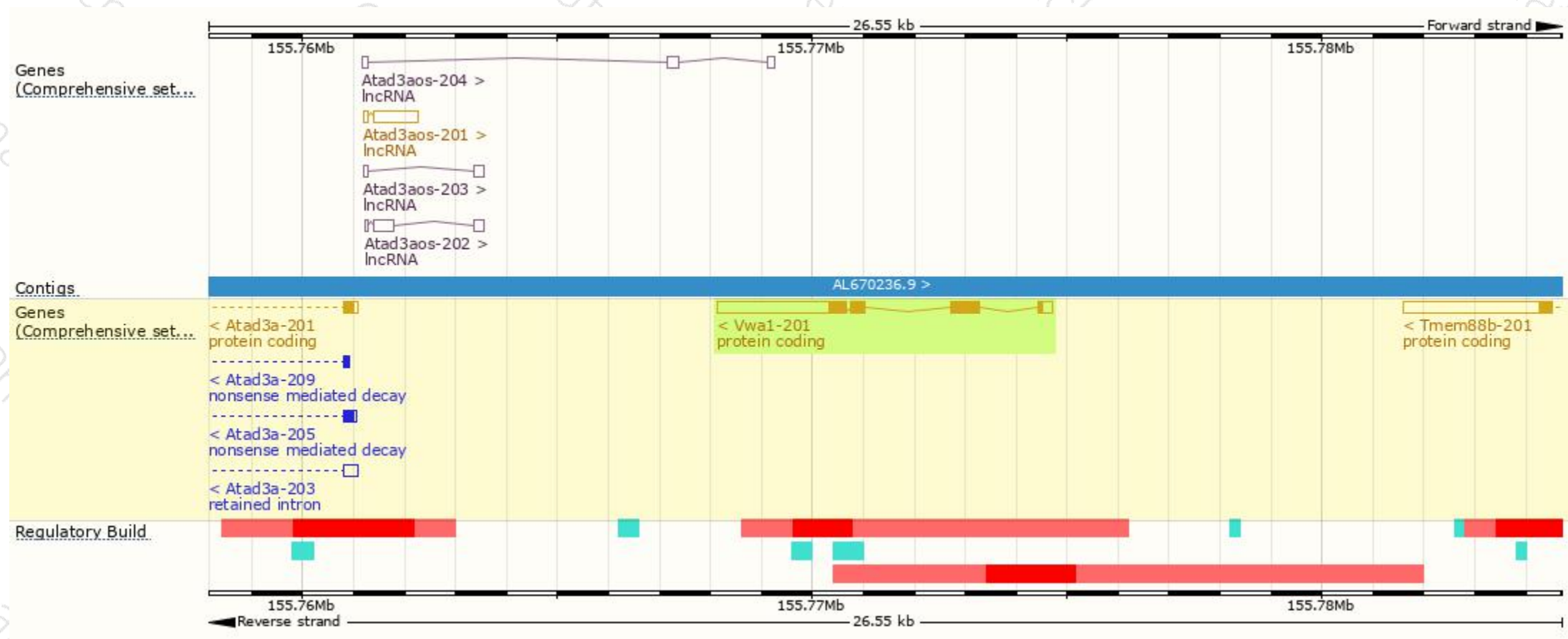
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Vwa1-201	ENSMUST00000042196.3	3623	415aa	Protein coding	CCDS19039	Q8R2Z5	TSL:1 GENCODE basic APPRIS P1

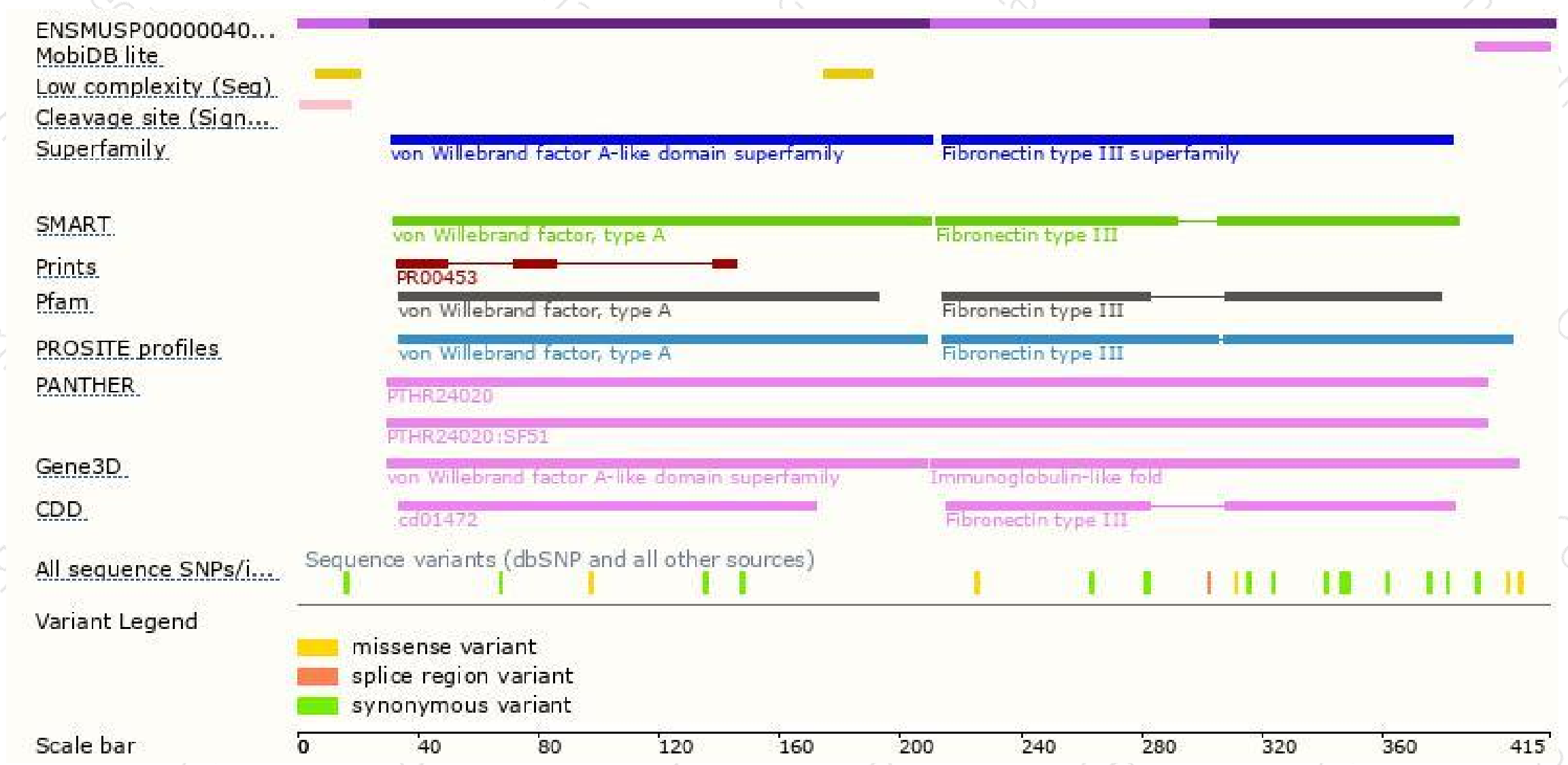
The strategy is based on the design of *Vwa1-201* transcript, The transcription is shown below



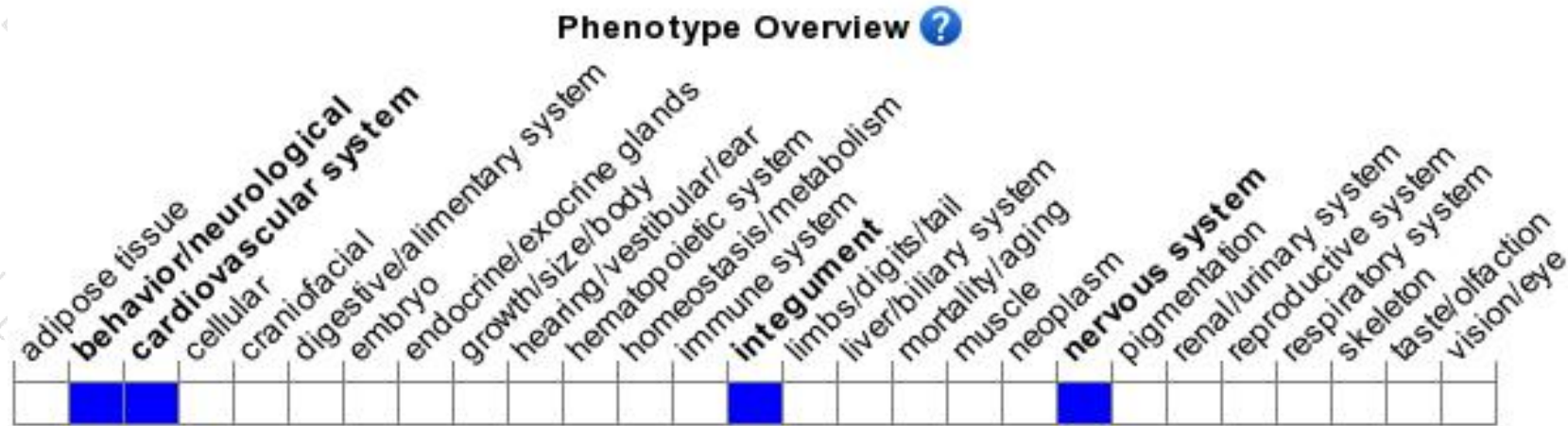
Genomic location distribution



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/>).

According to the existing MGI data, Mice homozygous for a reporter allele exhibit abnormal motor coordination/balance, increased thermal nociceptive threshold, and altered peripheral nerve structure and function.

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

Tel: 400-9660890

