

Igdcc3 Cas9-CKO Strategy

Designer: Xueting Zhang
reviewer: Yanhua Shen
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Project Overview

Project Name

Igdcc3

Project type

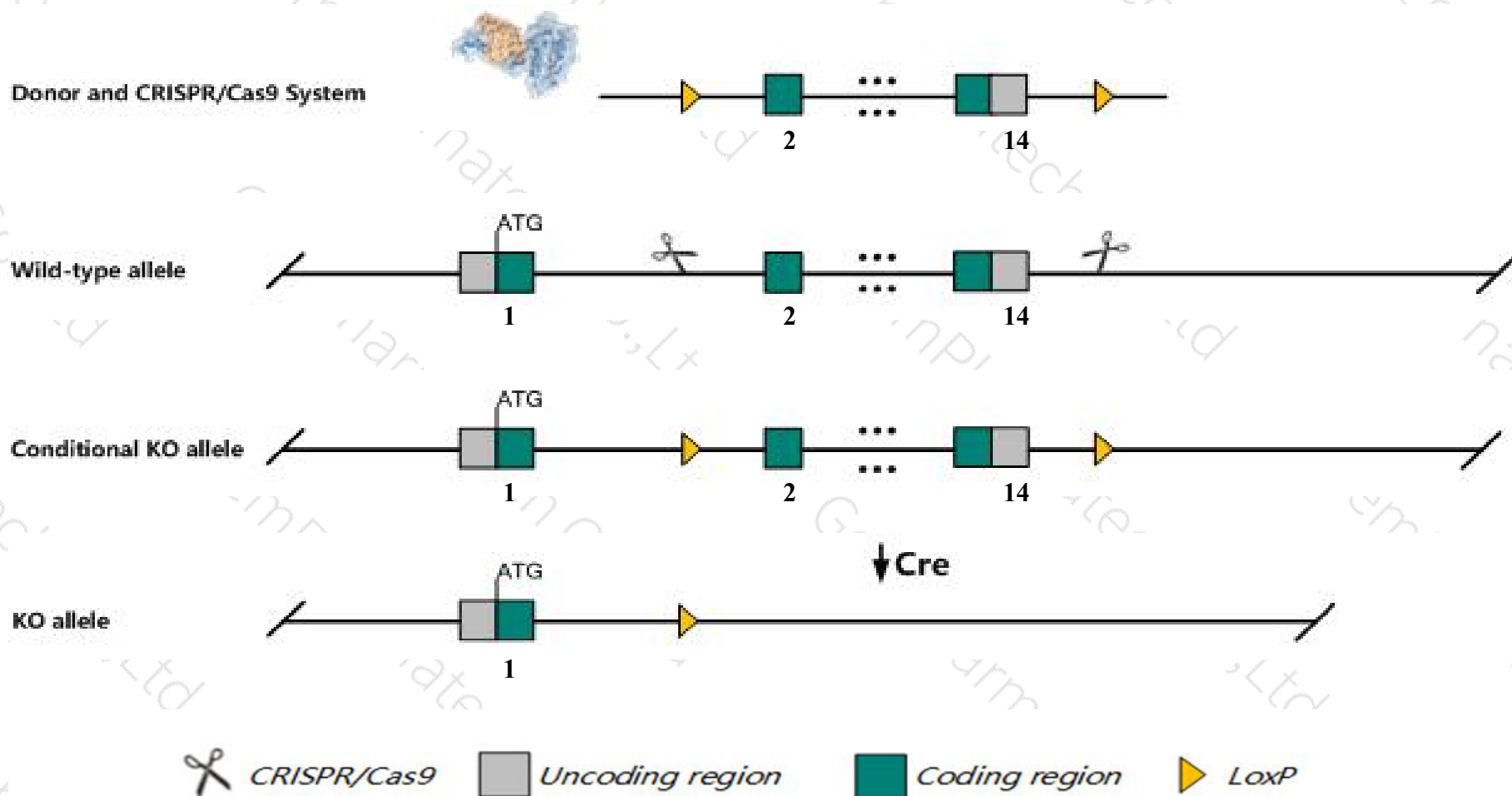
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Igdcc3* gene. The schematic diagram is as follows:



- The *Igdcc3* gene has 3 transcripts. According to the structure of *Igdcc3* gene, exon2-exon14 of *Igdcc3-201* (ENSMUST00000034961.5) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Igdcc3* 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, Homozygotes for a gene trap mutation exhibit reduced performance on the rotarod, suggesting impaired cerebellar function.
- The *Igdcc3* gene is located on the Chr9. 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)

Igdc3 immunoglobulin superfamily, DCC subclass, member 3 [*Mus musculus* (house mouse)]

Gene ID: 19289, updated on 24-Oct-2019

Summary

- Official Symbol

Igdc3 provided by MGI
- Official Full Name

immunoglobulin superfamily, DCC subclass, member 3 provided by MGI
- Primary source

[MGI:MGI:1202390](#)
- See related

[Ensembl:ENSMUSG00000032394](#)
- 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

Punc; AI851425; WI-14920; 2810401C09Rik
- Expression

Biased expression in CNS E11.5 (RPKM 19.1), CNS E14 (RPKM 4.0) and 5 other tissues [See more](#)
- Orthologs

[human](#) [all](#)

Genomic context

Location: 9 C; 9 35.13 cM

See Igdc3 in [Genome Data Viewer](#)

Exon count: 14

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	9	NC_000075.6 (65141154..65187031)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	9	NC_000075.5 (64988996..65033679)

Transcript information (Ensembl)

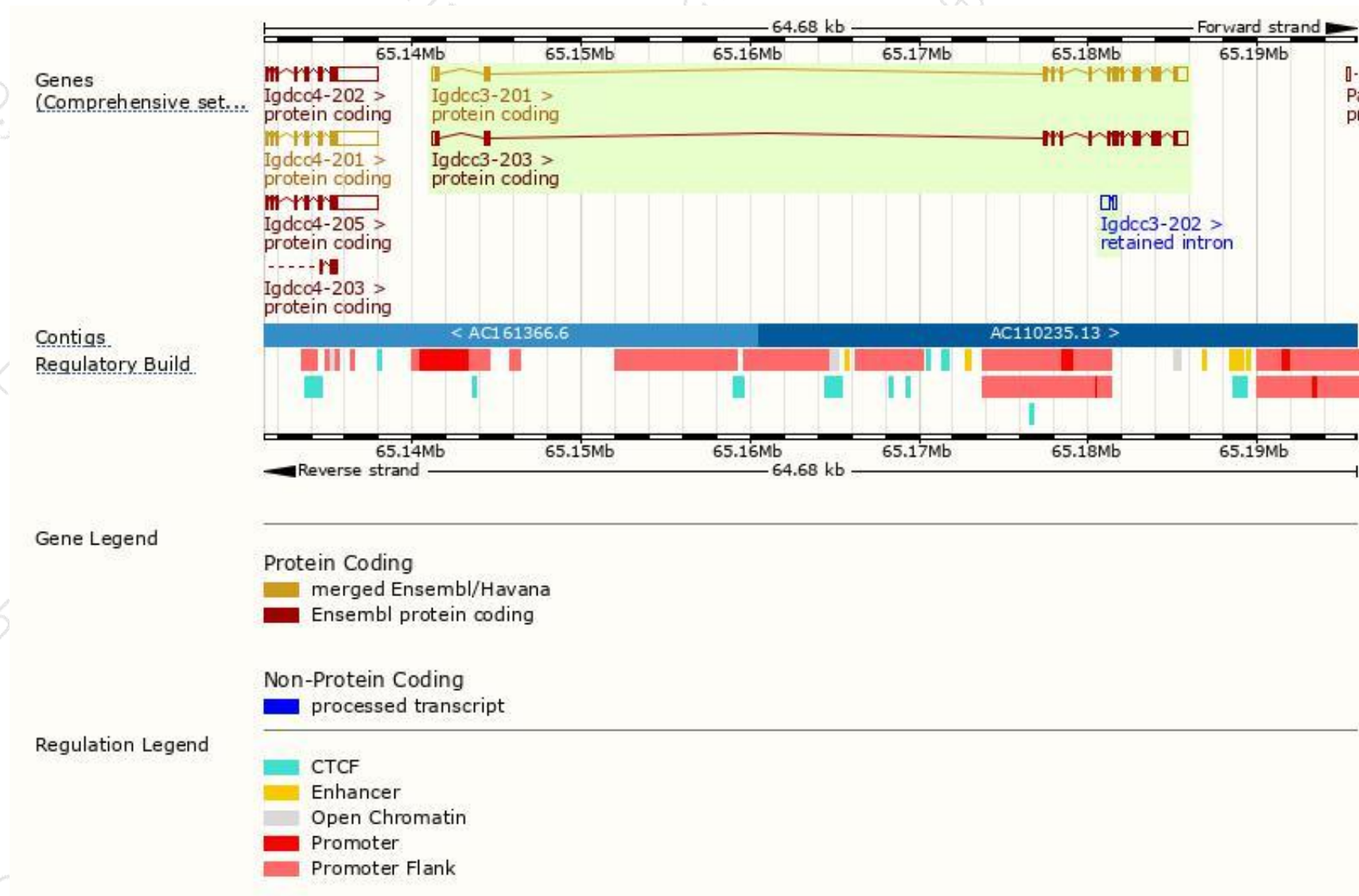
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Igdcc3-201	ENSMUST00000034961.5	3146	793aa	Protein coding	CCDS52835	Q8BQC3	TSL:1 GENCODE basic APPRIS P2
Igdcc3-203	ENSMUST00000217371.1	3183	813aa	Protein coding	-	Q8BQC3	TSL:1 GENCODE basic APPRIS ALT2
Igdcc3-202	ENSMUST00000217135.1	762	No protein	Retained intron	-	-	TSL:2

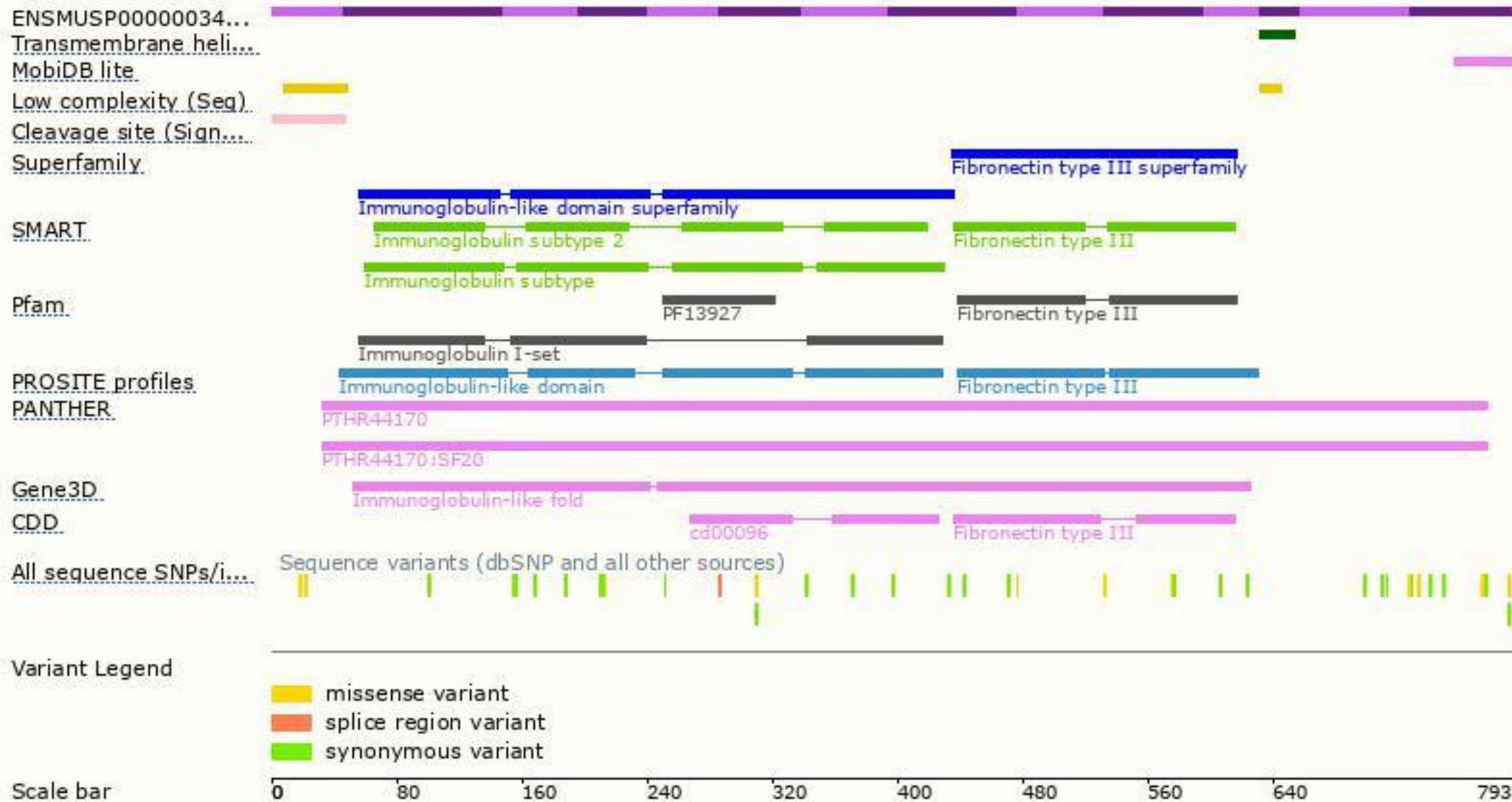
The strategy is based on the design of *Igdcc3-201* transcript,The transcription is shown below



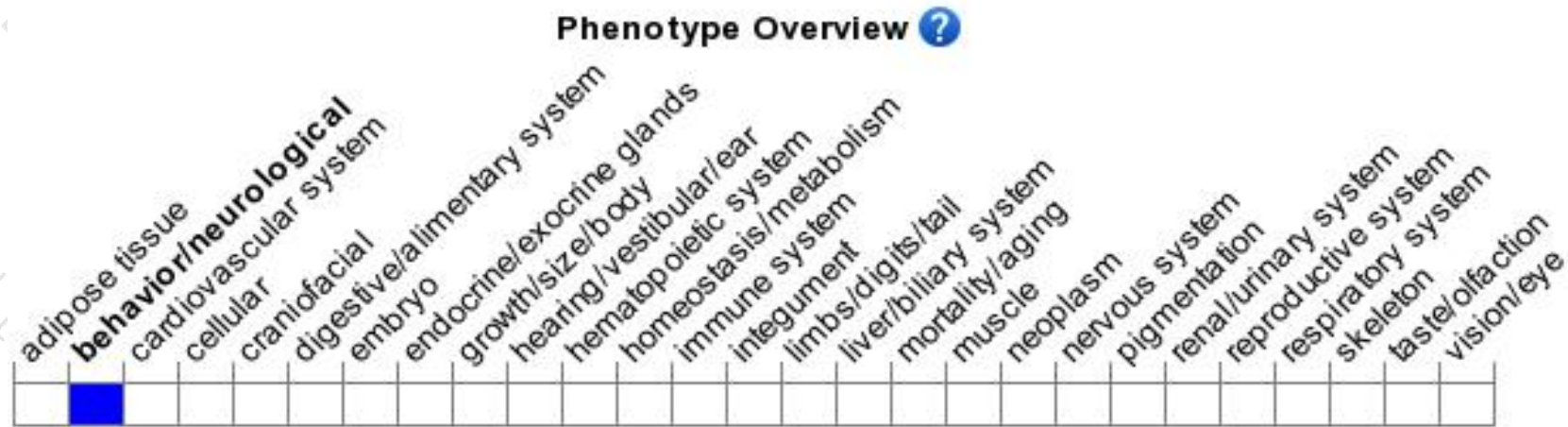
Genomic location distribution



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, Homozygotes for a gene trap mutation exhibit reduced performance on the rotarod, suggesting impaired cerebellar function.

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

Tel: 400-9660890

