

Gucy2f Cas9-KO Strategy

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Reviewer:

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Design Date:

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Project Overview

Project Name

Gucy2f

Project type

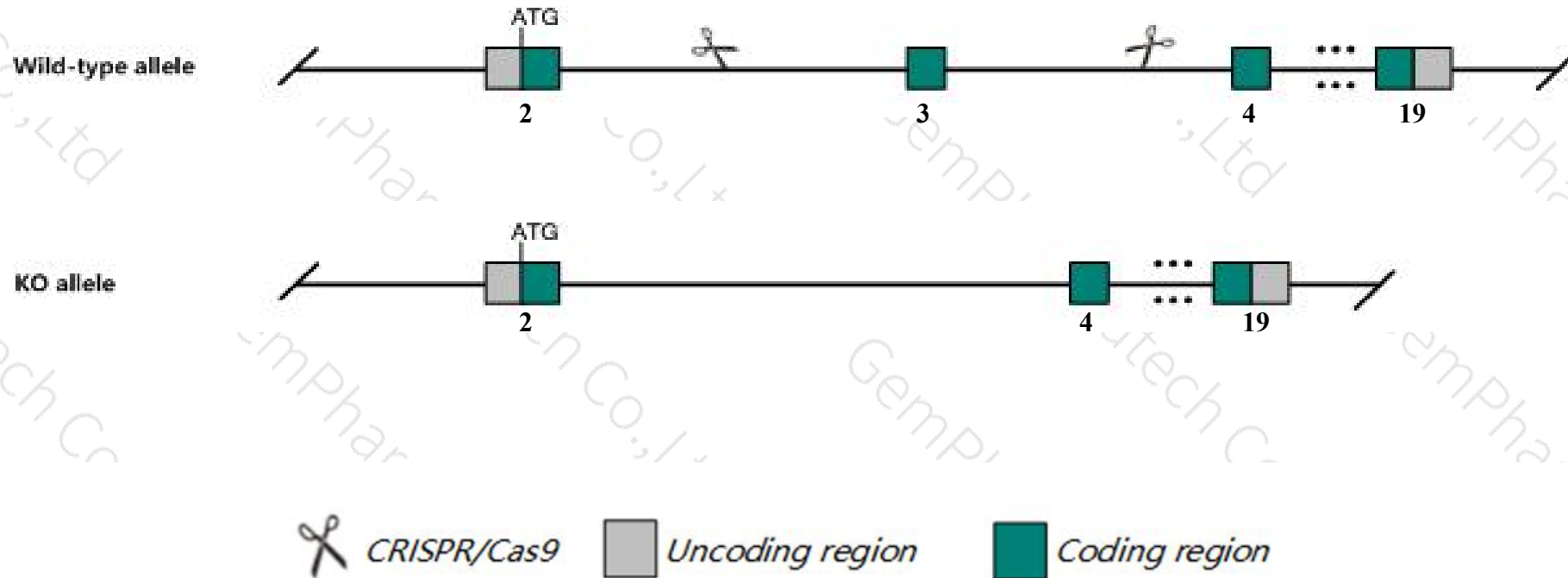
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gucy2f* gene. The schematic diagram is as follows:



- The *Gucy2f* gene has 1 transcript. According to the structure of *Gucy2f* gene, exon3 of *Gucy2f-201* (ENSMUST00000042530.3) transcript is recommended as the knockout region. The region contains 302bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gucy2f* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit abnormal retinal rod electrophysiology.
- The *Gucy2f* gene is located on the ChrX. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Gucy2f guanylate cyclase 2f [Mus musculus (house mouse)]

Gene ID: 245650, updated on 28-Mar-2019

Summary



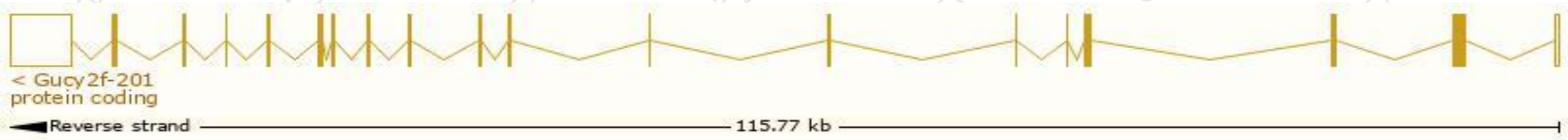
Official Symbol	Gucy2f provided by MGI
Official Full Name	guanylate cyclase 2f provided by MGI
Primary source	MGI:MGI:105119
See related	Ensembl:ENSMUSG00000042282
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	A930002I06, GC-F, GC2
Expression	Low expression observed in reference dataset See more
Orthologs	human all

Transcript information (Ensembl)

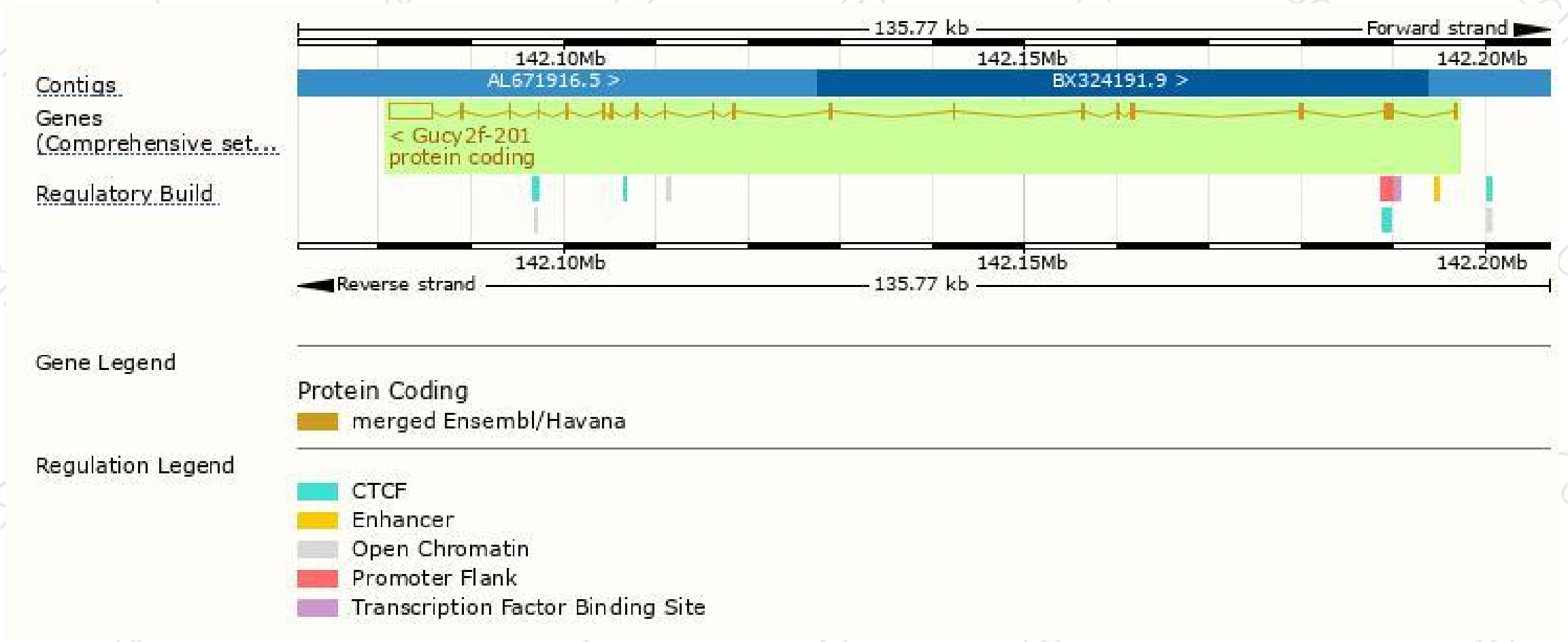
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gucy2f-201	ENSMUST00000042530.3	8122	1108aa	Protein coding	CCDS30446	Q5SDA5	TSL:1 GENCODE basic APPRIS P1

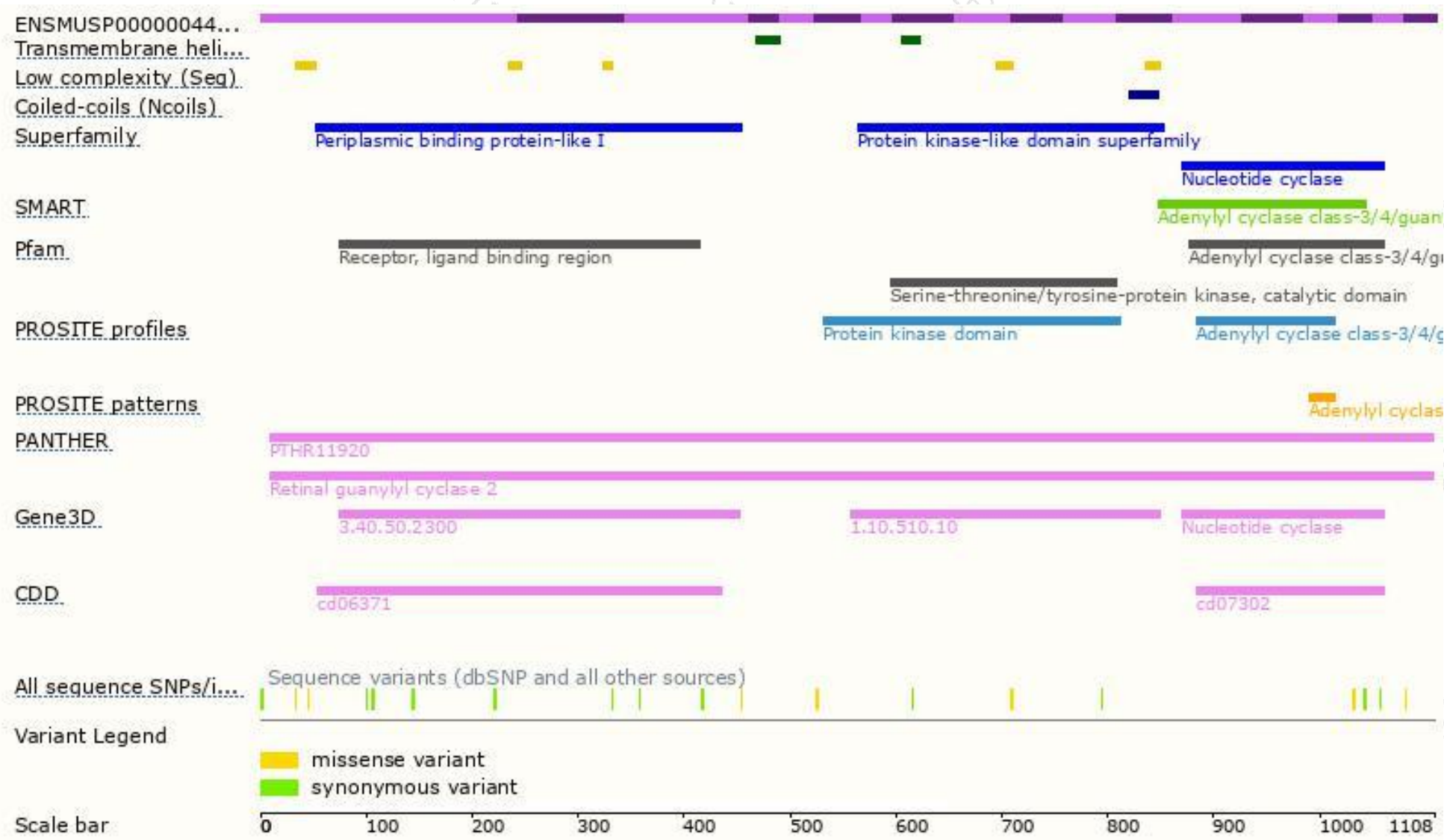
The strategy is based on the design of *Gucy2f-201* transcript, The transcription is shown below



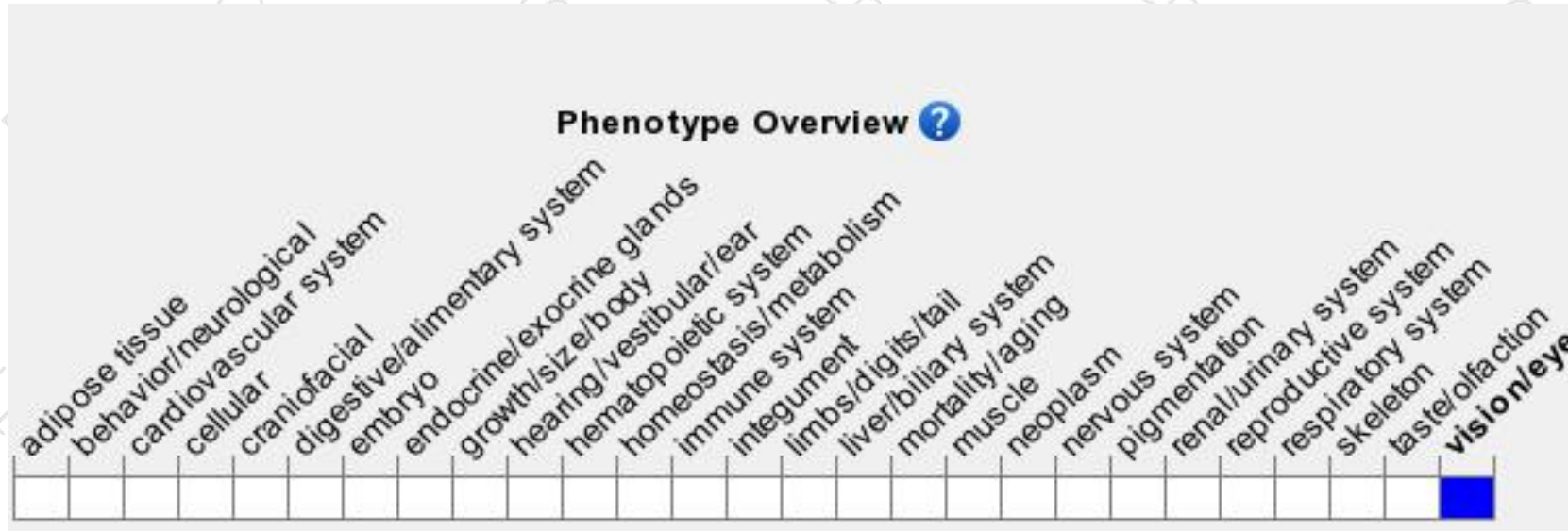
Genomic location distribution



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 knock-out allele exhibit abnormal retinal rod electrophysiology.

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

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