

Aldh3a1 Cas9-KO Strategy

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Project Overview

Project Name

Aldh3a1

Project type

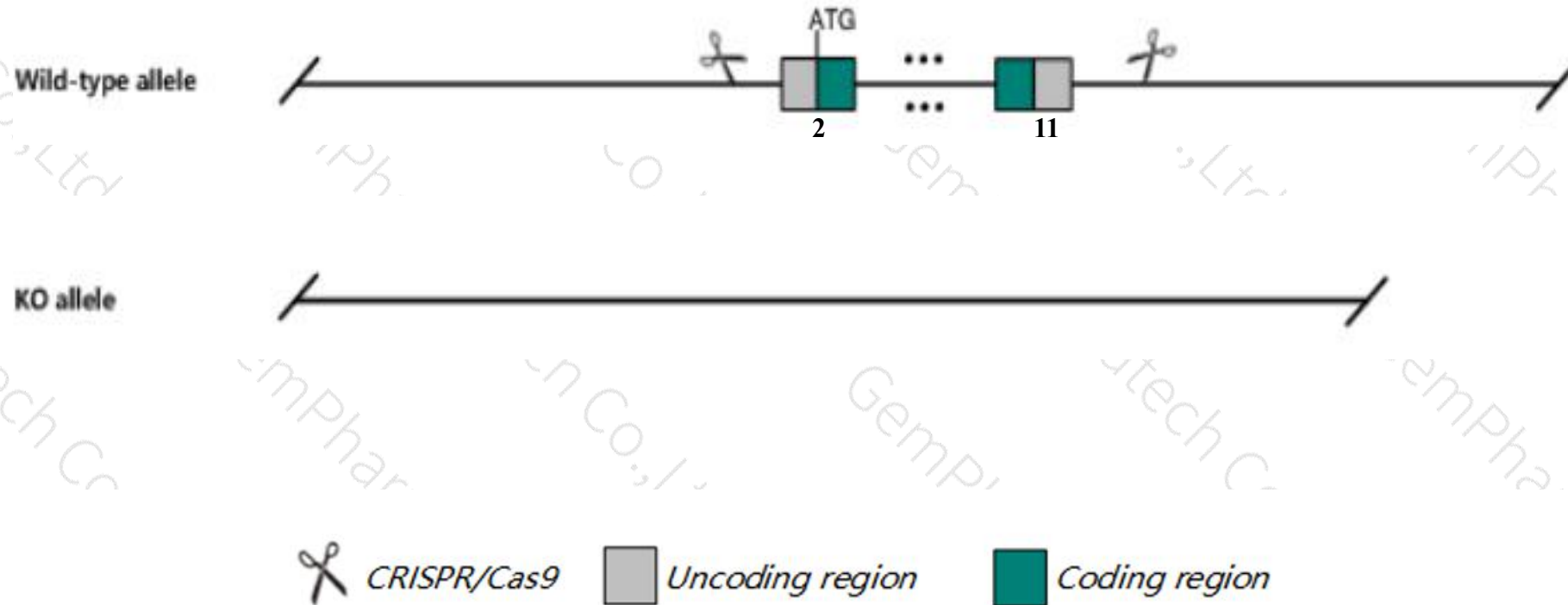
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Aldh3a1* gene. The schematic diagram is as follows:



- The *Aldh3a1* gene has 3 transcripts. According to the structure of *Aldh3a1* gene, exon2-exon11 of *Aldh3a1-201* (ENSMUST00000019246.3) transcript is recommended as the knockout region. The region contains all 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 *Aldh3a1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, homozygous mutation of this gene does not result in a phenotype.
- The *Aldh3a1* gene is located on the Chr11. 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)

Aldh3a1 aldehyde dehydrogenase family 3, subfamily A1 [Mus musculus (house mouse)]

Gene ID: 11670, updated on 13-Mar-2020

Summary



Official Symbol Aldh3a1 provided by [MGI](#)

Official Full Name aldehyde dehydrogenase family 3, subfamily A1 provided by [MGI](#)

Primary source [MGI:MGI:1353451](#)

See related [Ensembl:ENSMUSG00000019102](#)

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 Ahd-4, Ahd4, Aldh, Aldh3

Expression Biased expression in bladder adult (RPKM 55.3), stomach adult (RPKM 53.6) and 1 other tissue [See more](#)

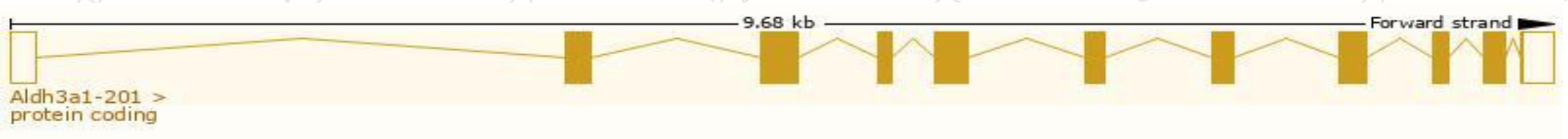
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

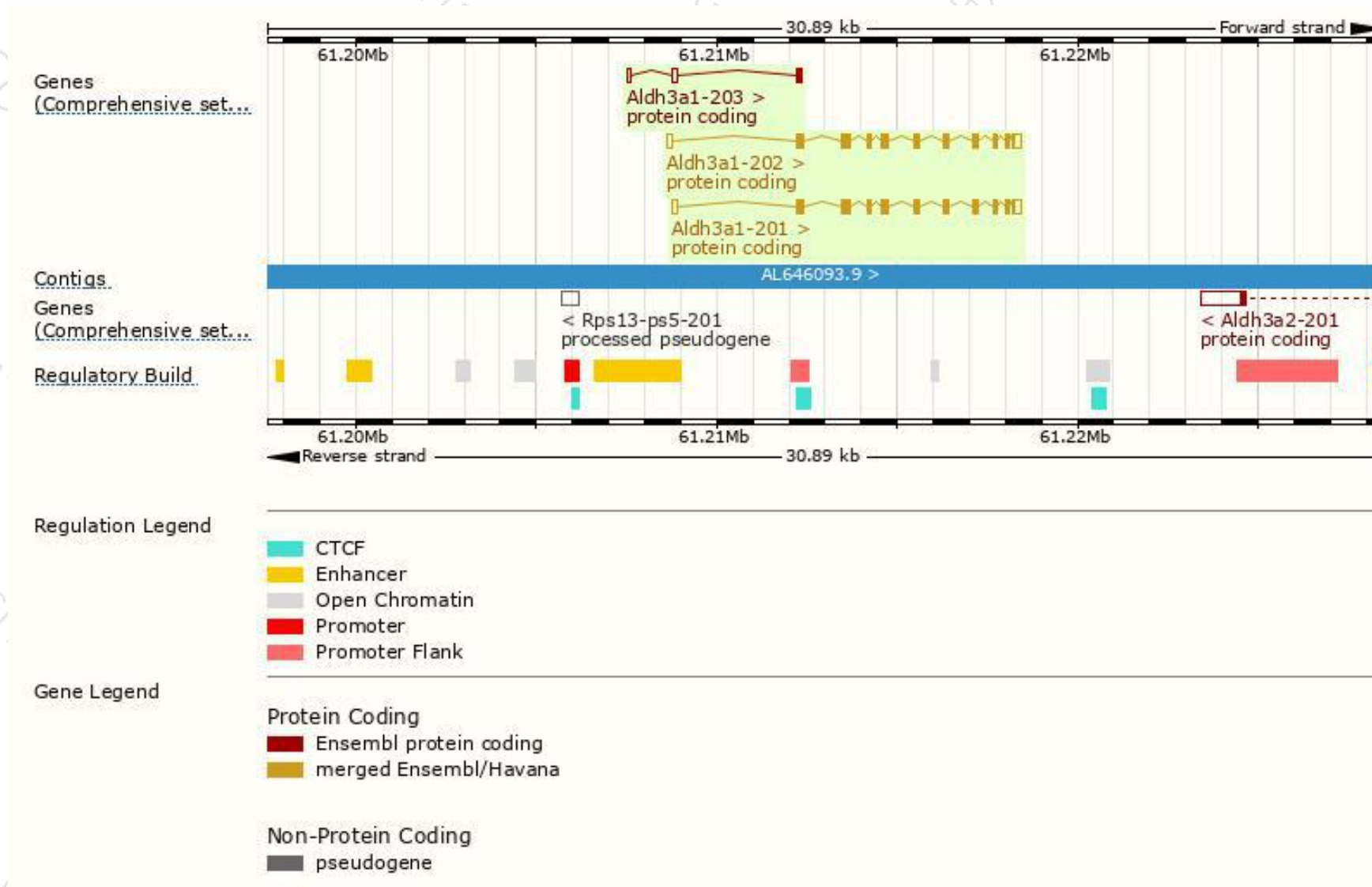
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Aldh3a1-201	ENSMUST00000019246.3	1729	453aa	Protein coding	CCDS24808	P47739	TSL:1 GENCODE basic APPRIS P1
Aldh3a1-202	ENSMUST00000108716.7	1721	453aa	Protein coding	CCDS24808	P47739	TSL:1 GENCODE basic APPRIS P1
Aldh3a1-203	ENSMUST00000147792.1	361	34aa	Protein coding	-	B1ATI6	CDS 3' incomplete TSL:3

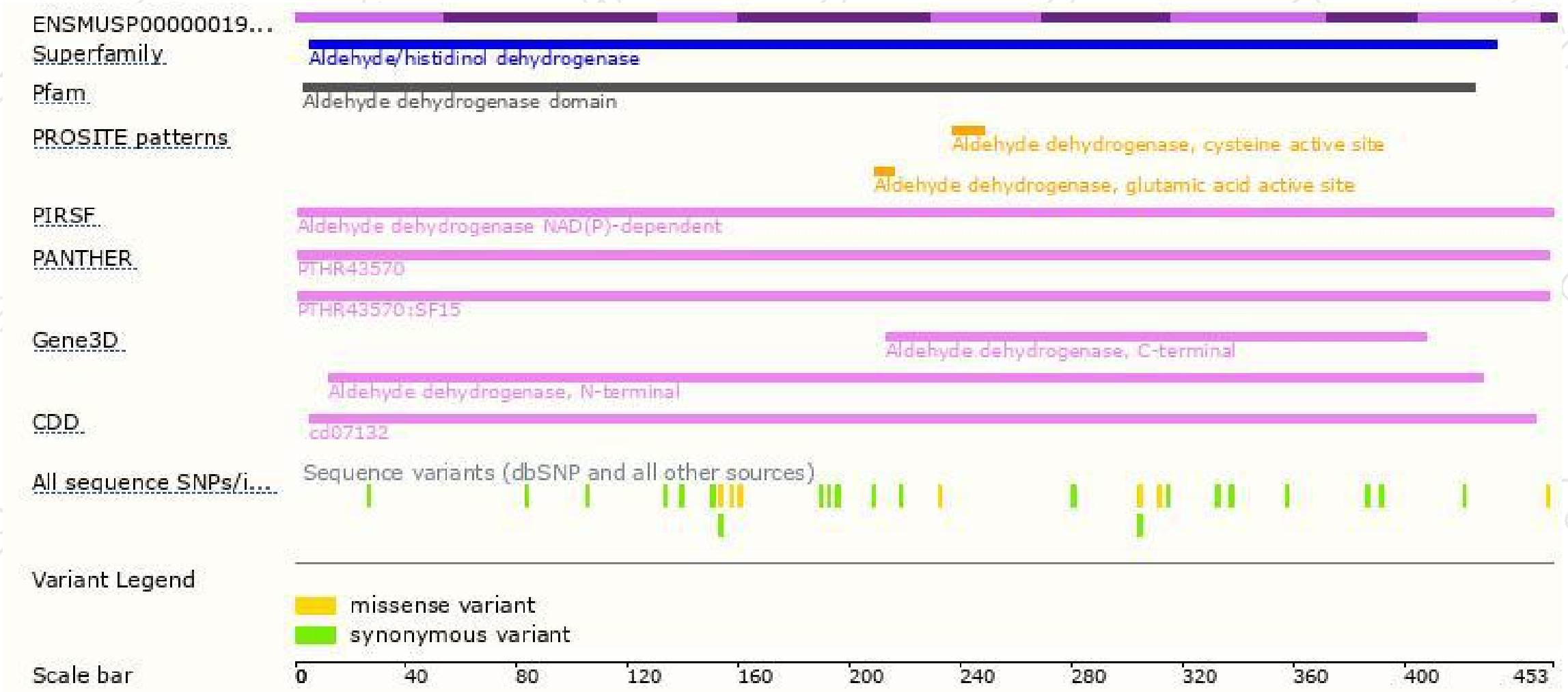
The strategy is based on the design of *Aldh3a1-201* transcript,the transcription is shown below:



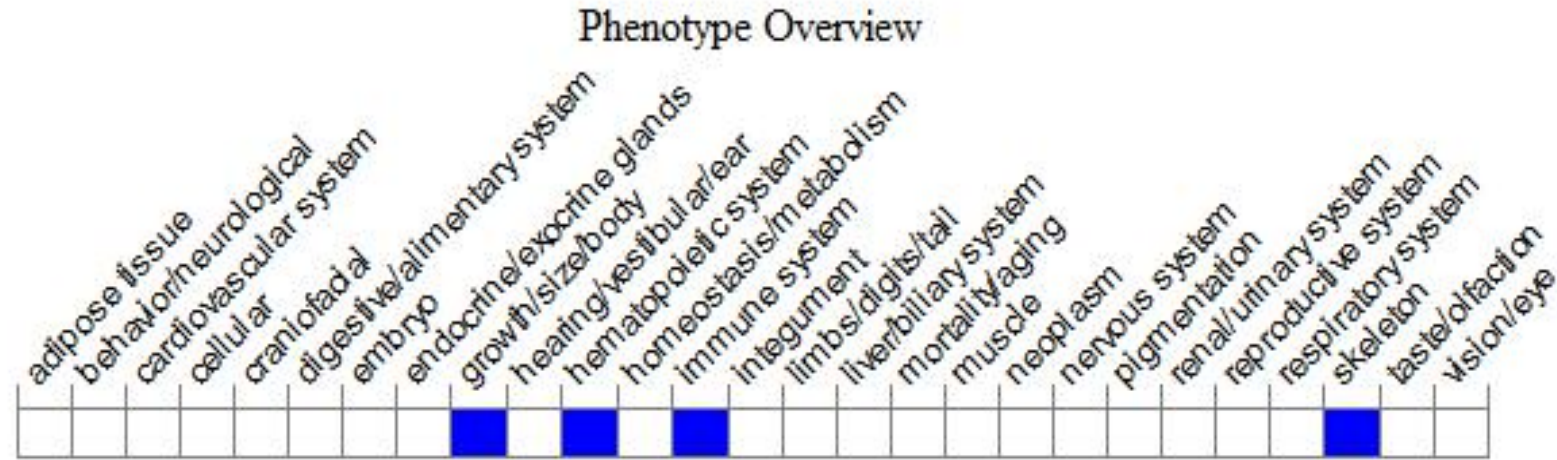
Genomic location distribution



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, homozygous mutation of this gene does not result in a phenotype.

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

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