

***Slitrk6* Cas9-KO Strategy**

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Project Overview

Project Name

Slitrk6

Project type

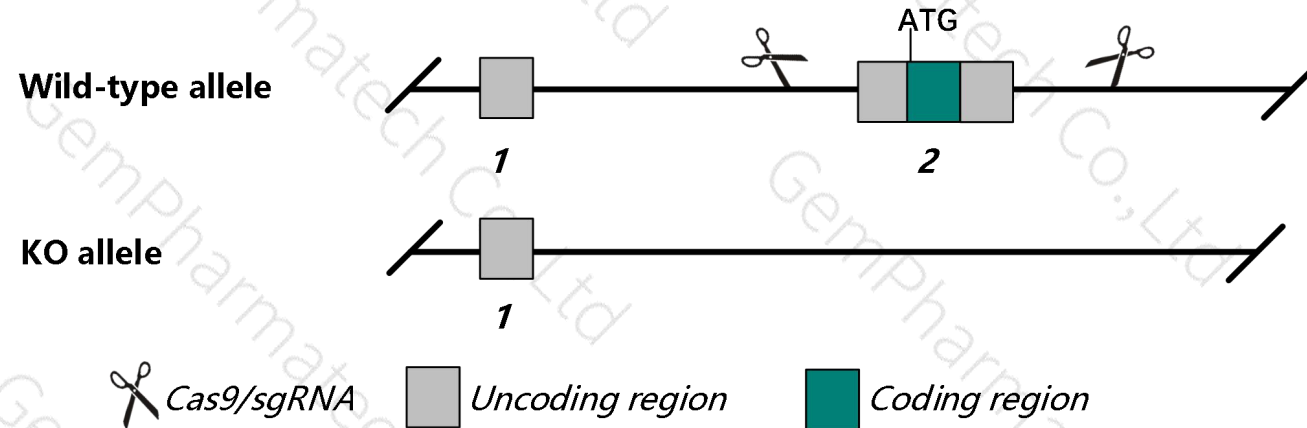
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slitrk6* gene. The schematic diagram is as follows:



- The *Slitrk6* gene has 1 transcript. According to the structure of *Slitrk6* gene, exon2 of *Slitrk6-201* (ENSMUST00000078386.3) 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 *Slitrk6* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous deficient mice show pronounced reduction in cochlear innervation. Innervation to the posterior crista is variably impaired and there is a loss of neurons in the spiral and vestibular ganglia.
- The *Slitrk6* gene is located on the Chr14. 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)

Slitrk6 SLIT and NTRK-like family, member 6 [*Mus musculus* (house mouse)]

Gene ID: 239250, updated on 24-Dec-2019

Summary

- Official Symbol** Slitrk6 provided by MGI
- Official Full Name** SLIT and NTRK-like family, member 6 provided by MGI
- Primary source** MGI:MGI:2443198
- See related** Ensembl:ENSMUSG00000045871
- 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** Sltk6; 4832410J21Rik
- Expression** Biased expression in bladder adult (RPKM 3.5), limb E14.5 (RPKM 3.4) and 10 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: 14; 14 E3 See Slitrk6 in [Genome Data Viewer](#)

Exon count: 2

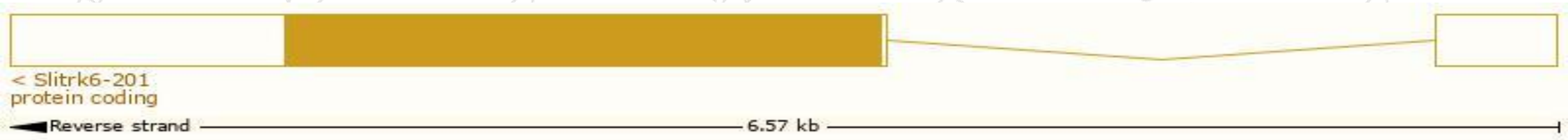
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	14	NC_000080.6 (110747338..110755204, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	14	NC_000080.5 (111147800..111154371, complement)

Transcript information (Ensembl)

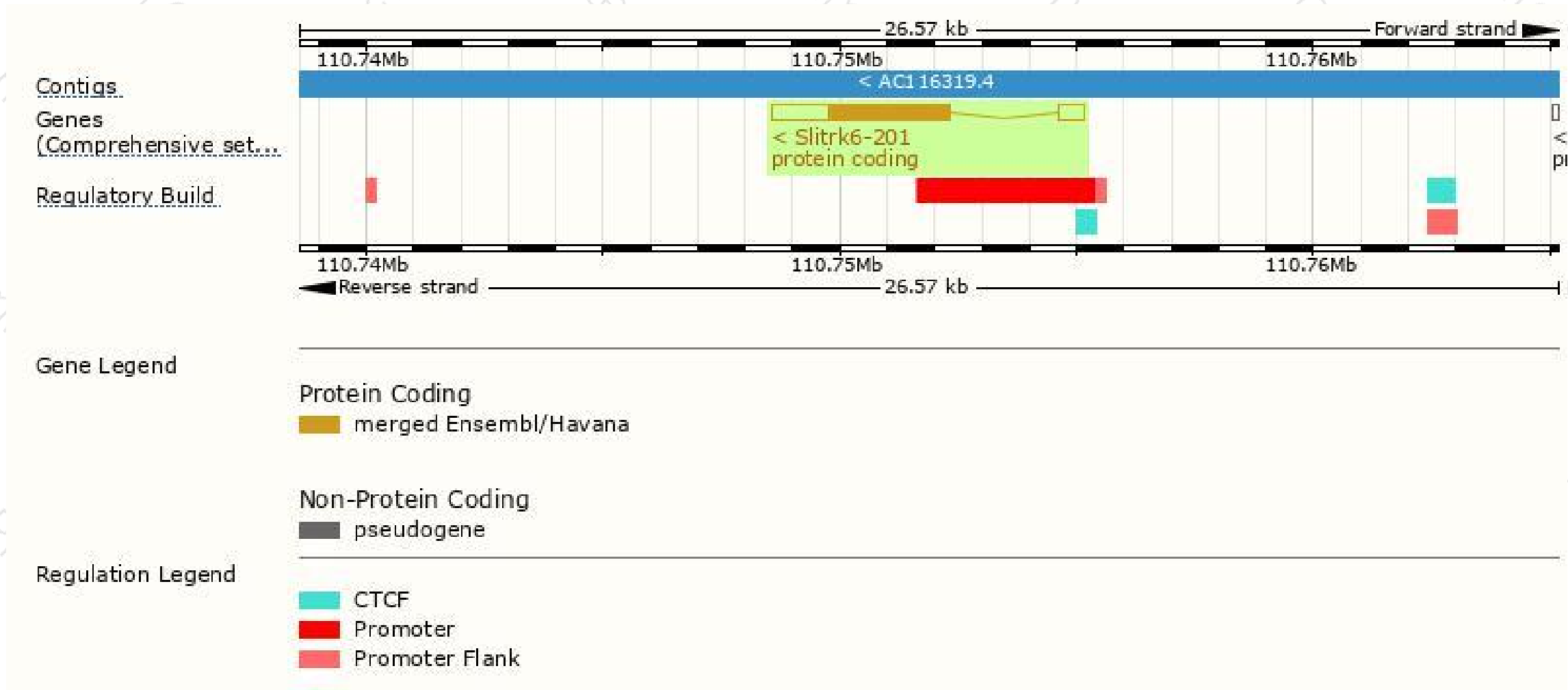
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slitrk6-201	ENSMUST00000078386.3	4240	840aa	Protein coding	CCDS27326	A6H6M2 Q8C110	TSL:1 GENCODE basic APPRIS P1

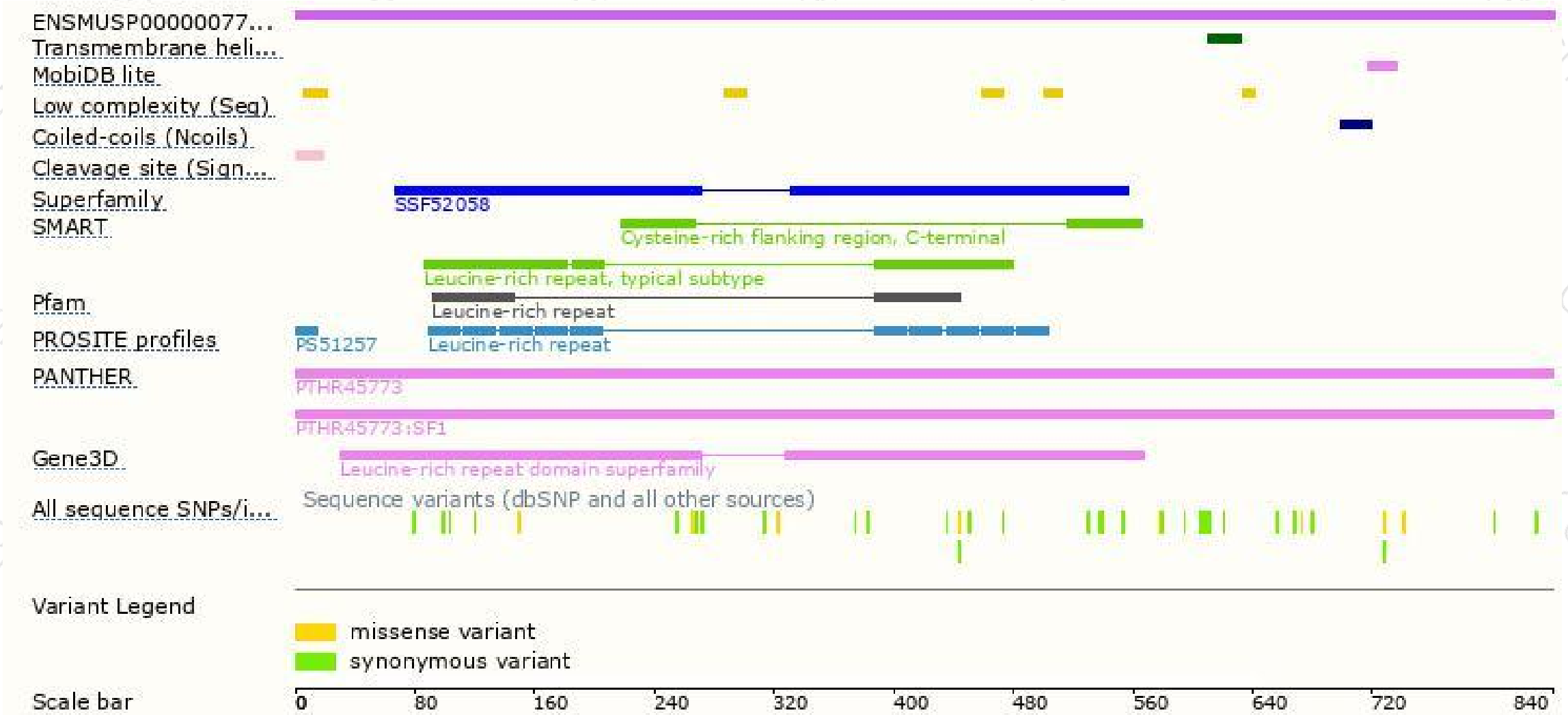
The strategy is based on the design of *Slitrk6-201* transcript,The transcription is shown below



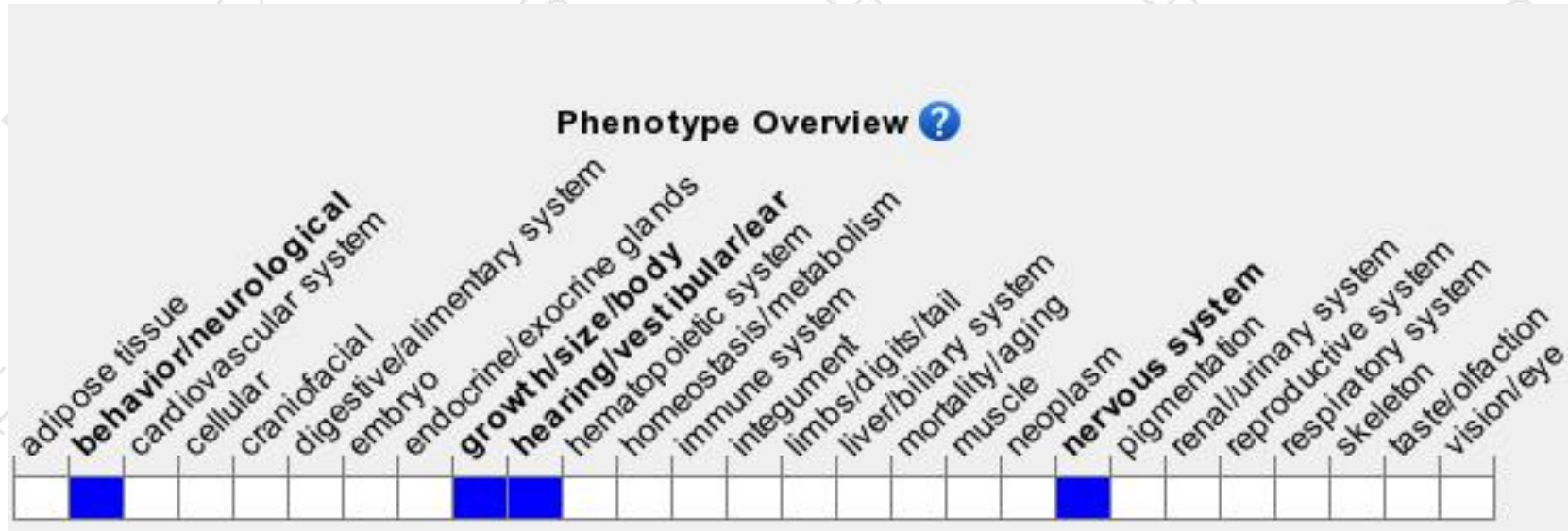
Genomic location distribution



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 deficient mice show pronounced reduction in cochlear innervation.

Innervation to the posterior crista is variably impaired and there is a loss of neurons in the spiral and vestibular ganglia.

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

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