

# *Grxcr2* Cas9-KO Strategy

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Reviewer Xueting Zhang  
Design Date: 2020-3-11

# Project Overview

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|---------------------|----------------------|
| <b>Project Name</b> | <b><i>Grxcr2</i></b> |
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|---------------------|----------------|
| <b>Project type</b> | <b>Cas9-KO</b> |
|---------------------|----------------|

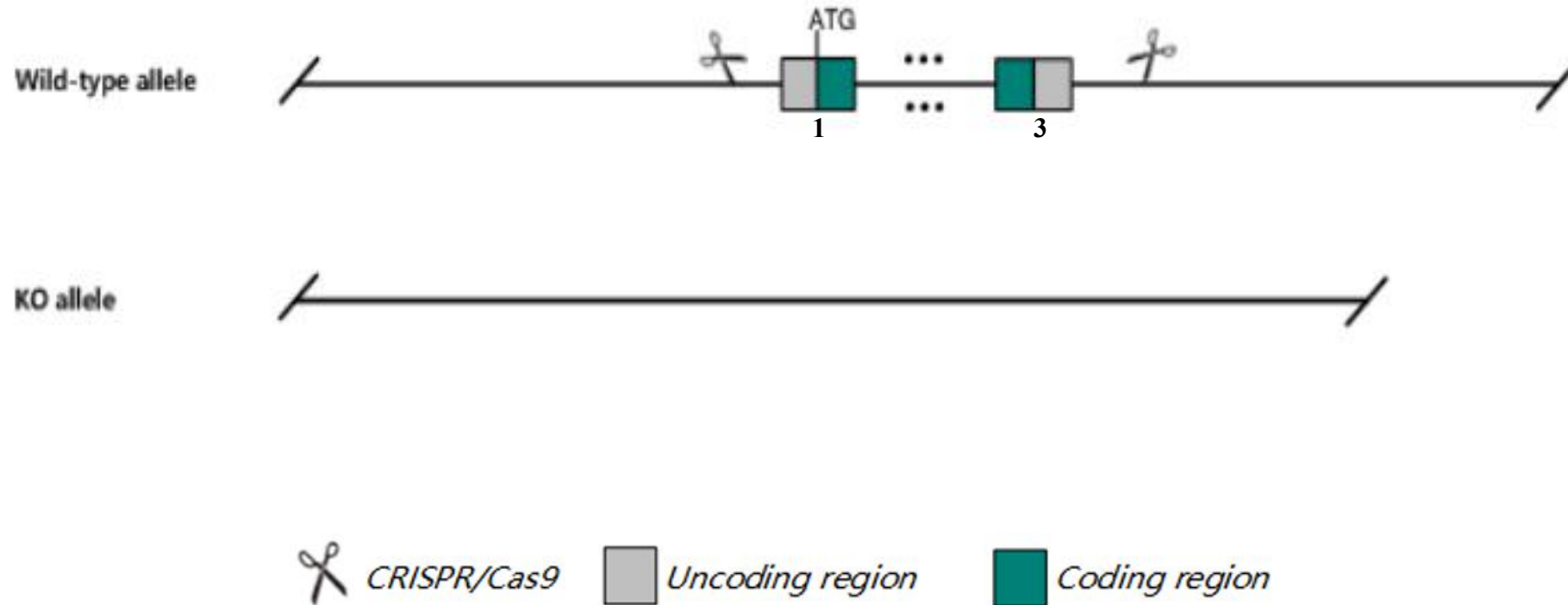
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|--------------------------|--------------------|
| <b>Strain background</b> | <b>C57BL/6JGpt</b> |
|--------------------------|--------------------|

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Grxcr2* gene. The schematic diagram is as follows:



The *Grxcr2* gene has 1 transcript. According to the structure of *Grxcr2* gene, exon1-exon3 of *Grxcr2-201* (ENSMUST00000097591.4) 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 *Grxcr2* gene. The brief process is as follows: CRISPR/Cas9 system

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit severe hearing loss.

The *Grxcr2* gene is located on the Chr18. 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.

**Grxcr2** glutaredoxin, cysteine rich 2 [ *Mus musculus* (house mouse) ]

Gene ID: 332309, updated on 27-Feb-2020

## Summary

|                    |                                                                                                                                                                           |
|--------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Official Symbol    | Grxcr2 provided by <a href="#">MGI</a>                                                                                                                                    |
| Official Full Name | glutaredoxin, cysteine rich 2 provided by <a href="#">MGI</a>                                                                                                             |
| Primary source     | <a href="#">MGI:MGI:2685697</a>                                                                                                                                           |
| See related        | <a href="#">Ensembl:ENSMUSG00000073574</a>                                                                                                                                |
| Gene type          | protein coding                                                                                                                                                            |
| RefSeq status      | VALIDATED                                                                                                                                                                 |
| Organism           | <a href="#">Mus musculus</a>                                                                                                                                              |
| Lineage            | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| Also known as      | Gm851                                                                                                                                                                     |
| Expression         | Low expression observed in reference dataset <a href="#">See more</a>                                                                                                     |
| Orthologs          | <a href="#">human</a> <a href="#">all</a>                                                                                                                                 |

## Genomic context

Location: 18; 18 B3

Exon count: 3

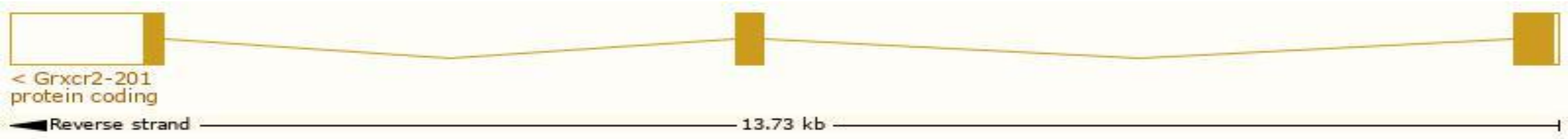
See Grxcr2 in [Genome Data Viewer](#)

# Transcript information      Ensembl

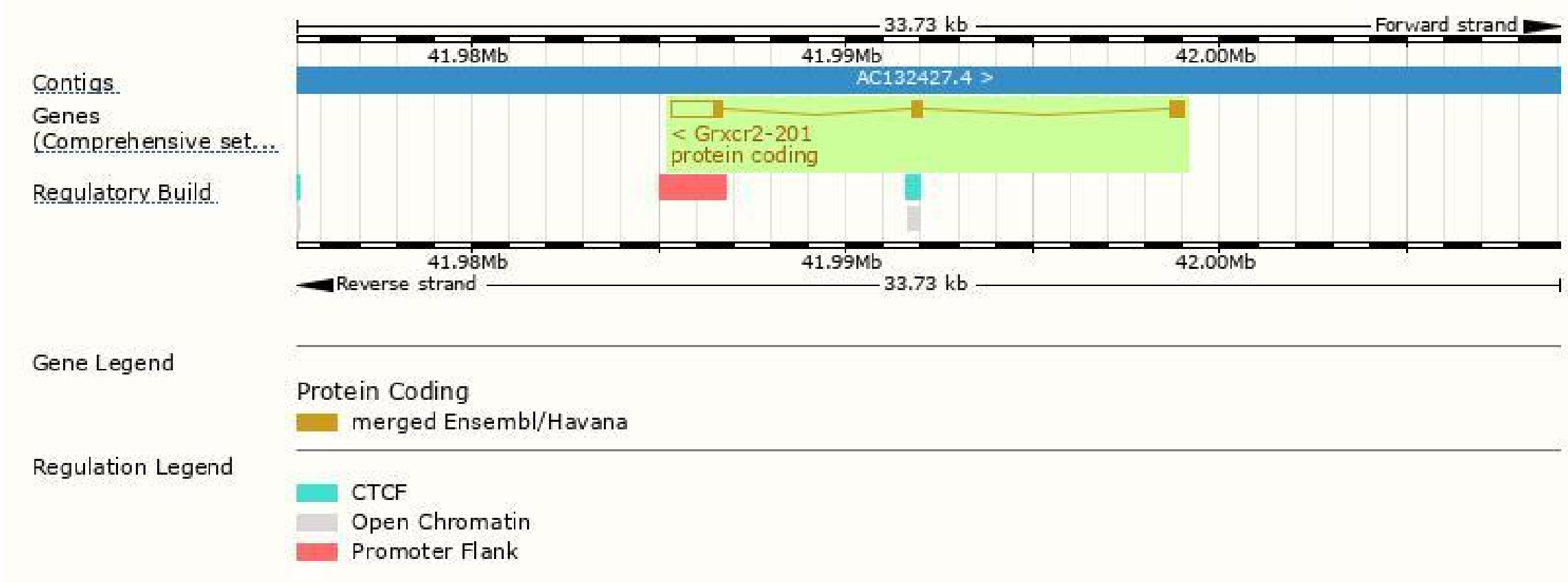
The gene has 1 transcript,and the transcript is shown below:

| Name       | Transcript ID                        | bp   | Protein               | Biotype        | CCDS                      | UniProt                | Flags                         |
|------------|--------------------------------------|------|-----------------------|----------------|---------------------------|------------------------|-------------------------------|
| Grxcr2-201 | <a href="#">ENSMUST00000097591.4</a> | 1998 | <a href="#">254aa</a> | Protein coding | <a href="#">CCDS29209</a> | <a href="#">Q3TYR5</a> | TSL:1 GENCODE basic APPRIS P1 |

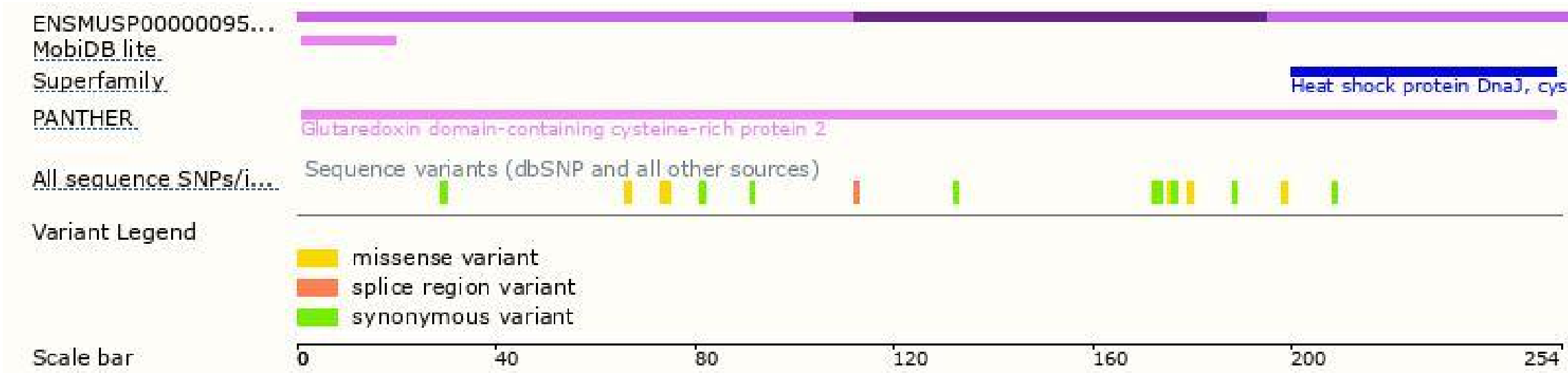
The strategy is based on the design of *Grxcr2-201* transcript,The transcription is shown below



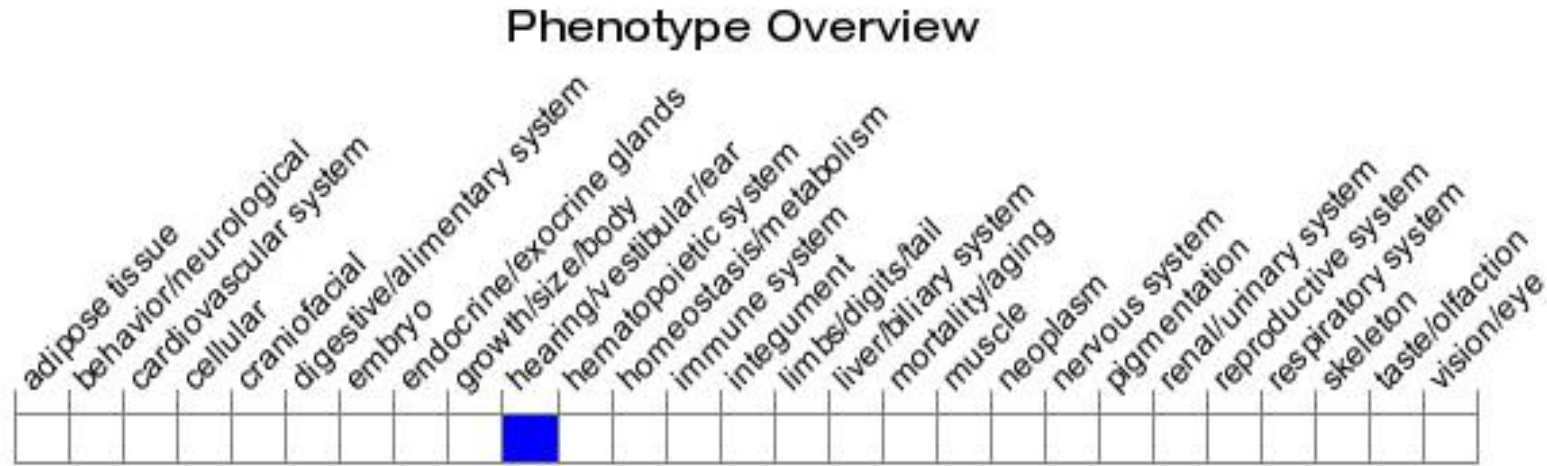
# Genomic location distribution



# 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 severe hearing loss.

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