

# ***Kirrel3 Cas9-KO Strategy***

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**Reviewer: Yumeng Wang**

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# Project Overview

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**Project Name**

***Kirrel3***

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**Project type**

**Cas9-KO**

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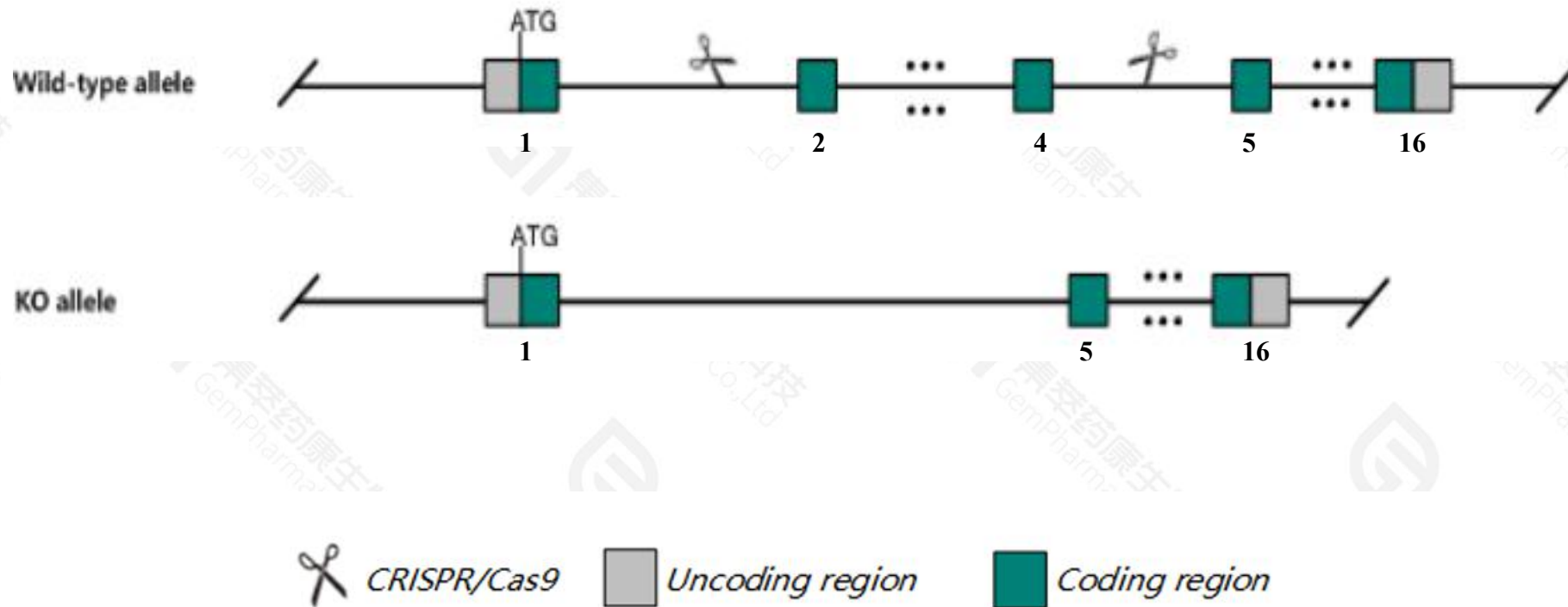
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Kirrel3* gene. The schematic diagram is as follows:



- The *Kirrel3* gene has 9 transcripts. According to the structure of *Kirrel3* gene, exon2-exon4 of *Kirrel3*-205(ENSMUST00000187182.7) transcript is recommended as the knockout region. The region contains 458bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Kirrel3* gene. The brief process is as follows: CRISPR/Cas9 system 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.



- According to the existing MGI data, mice homozygous for a knock-out allele exhibit impaired accessory olfactory bulb formation with reduced coalescence of vomeronasal sensory neuron axons in the posterior accessory olfactory bulb, loss of male-male aggression and abnormal male sexual response to a male intruder mouse.
- The *Kirrel3* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

## Kirrel3 kirre like nephrin family adhesion molecule 3 [Mus musculus (house mouse)]

Gene ID: 67703, updated on 17-Dec-2020

### Summary



**Official Symbol** Kirrel3 provided by [MGI](#)

**Official Full Name** kirre like nephrin family adhesion molecule 3 provided by [MGI](#)

**Primary source** [MGI:MGI:1914953](#)

**See related** [Ensembl:ENSMUSG00000032036](#)

**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** 1500010O20Rik, 2900036G11Rik, NEPH2, Nep, SST4, mKIAA1867

**Expression** Biased expression in cerebellum adult (RPKM 9.9), frontal lobe adult (RPKM 5.7) and 7 other tissues [See more](#)

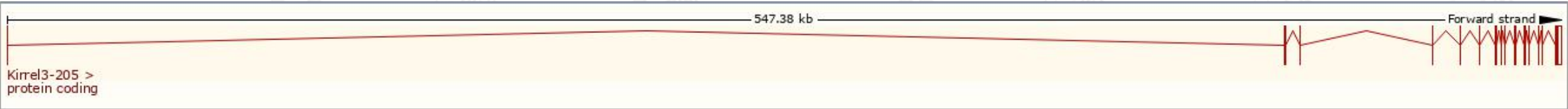
**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

The gene has 9 transcripts,all transcripts are shown below:

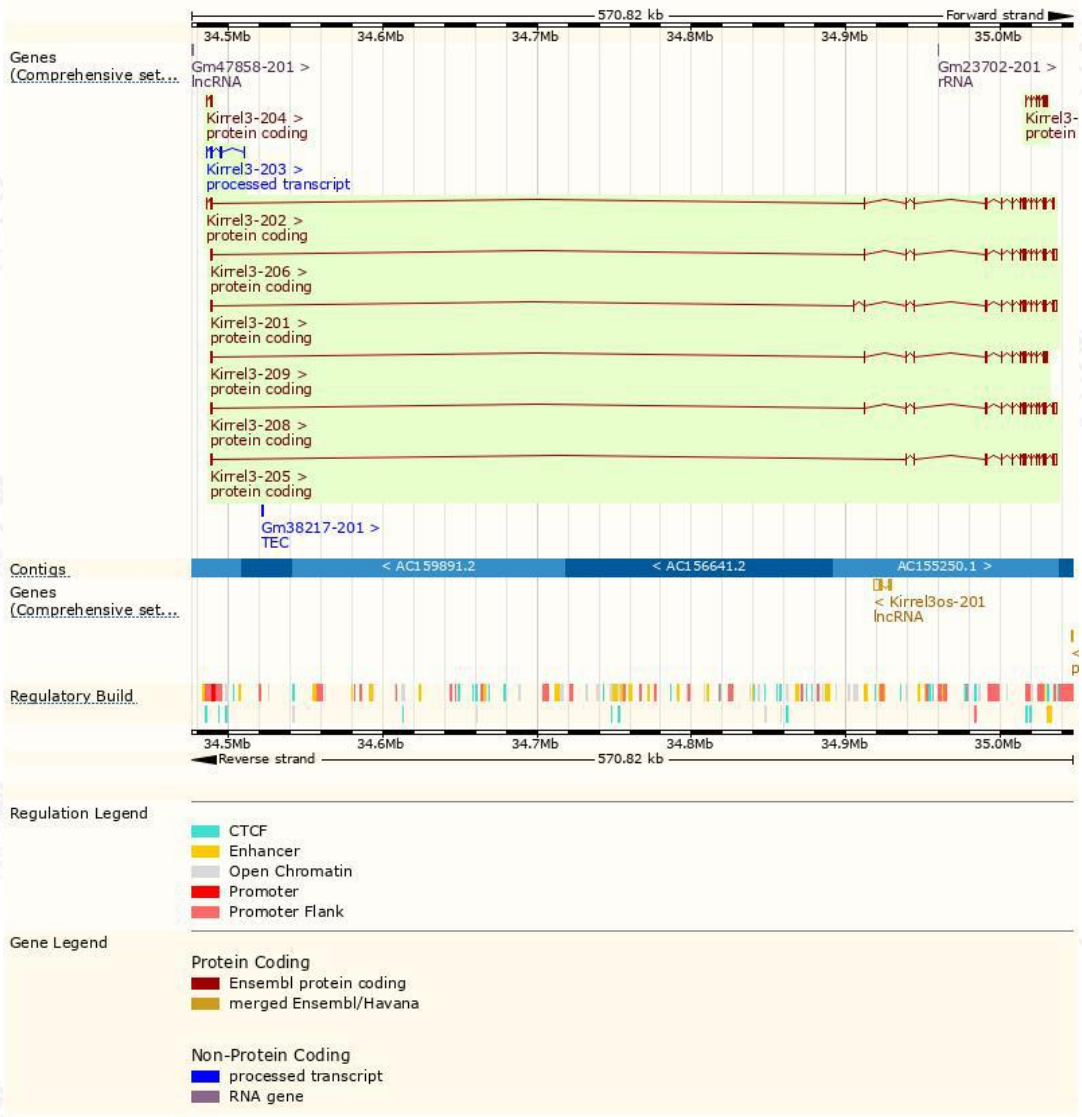
| Name        | Transcript ID                         | bp   | Protein               | Biotype              | CCDS                      | UniProt | Flags                               |
|-------------|---------------------------------------|------|-----------------------|----------------------|---------------------------|---------|-------------------------------------|
| Kirrel3-201 | <a href="#">ENSMUST00000045091.13</a> | 4147 | <a href="#">733aa</a> | Protein coding       | <a href="#">CCDS57667</a> |         | TSL:1 , GENCODE basic ,             |
| Kirrel3-206 | <a href="#">ENSMUST00000187625.7</a>  | 4108 | <a href="#">733aa</a> | Protein coding       | <a href="#">CCDS57667</a> |         | TSL:1 , GENCODE basic ,             |
| Kirrel3-205 | <a href="#">ENSMUST00000187182.7</a>  | 3532 | <a href="#">777aa</a> | Protein coding       | <a href="#">CCDS80973</a> |         | TSL:1 , GENCODE basic ,             |
| Kirrel3-202 | <a href="#">ENSMUST00000115148.9</a>  | 2851 | <a href="#">766aa</a> | Protein coding       | <a href="#">CCDS40574</a> |         | TSL:5 , GENCODE basic , APPRIS P1 , |
| Kirrel3-208 | <a href="#">ENSMUST00000188933.7</a>  | 3643 | <a href="#">745aa</a> | Protein coding       | -                         |         | TSL:1 , GENCODE basic ,             |
| Kirrel3-209 | <a href="#">ENSMUST00000190549.7</a>  | 3074 | <a href="#">579aa</a> | Protein coding       | -                         |         | TSL:1 , GENCODE basic ,             |
| Kirrel3-207 | <a href="#">ENSMUST00000188658.2</a>  | 1644 | <a href="#">239aa</a> | Protein coding       | -                         |         | CDS 5' incomplete , TSL:1 ,         |
| Kirrel3-204 | <a href="#">ENSMUST00000184203.8</a>  | 746  | <a href="#">18aa</a>  | Protein coding       | -                         |         | CDS 3' incomplete , TSL:3 ,         |
| Kirrel3-203 | <a href="#">ENSMUST00000183580.3</a>  | 1334 | No protein            | Processed transcript | -                         |         | TSL:1 ,                             |

The strategy is based on the design of *Kirrel3-205* transcript,the transcription is shown below:



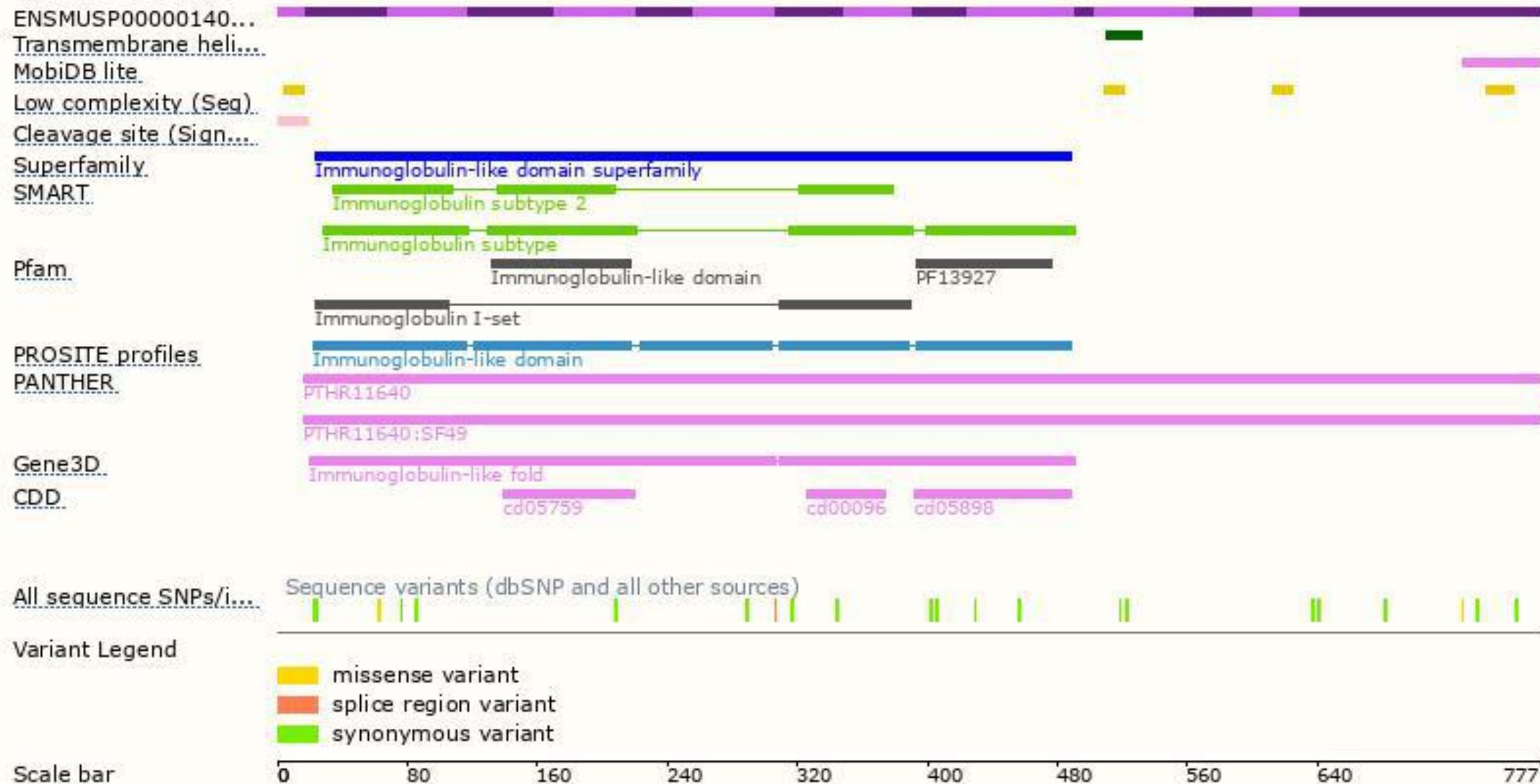


# Genomic location distribution

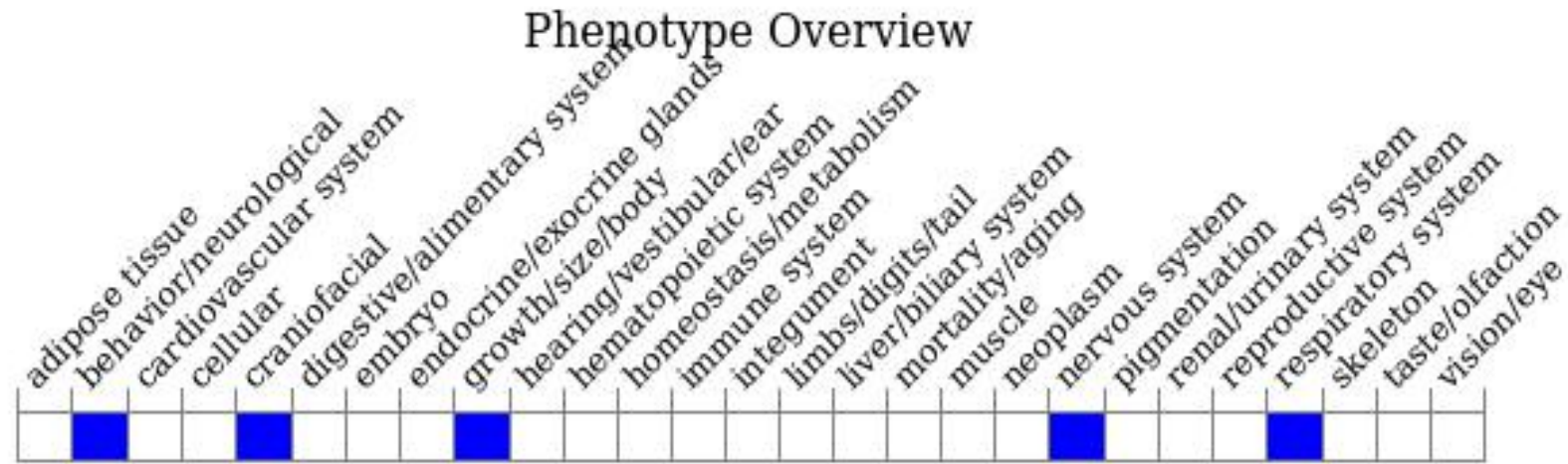




# Protein domain



# 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 impaired accessory olfactory bulb formation with reduced coalescence of vomeronasal sensory neuron axons in the posterior accessory olfactory bulb, loss of male-male aggression and abnormal male sexual response to a male intruder mouse.

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