

# ***Sohlh2* Cas9-KO Strategy**

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**Reviewer: Jia Yu**

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# Project Overview

**Project Name**

***Sohlh2***

**Project type**

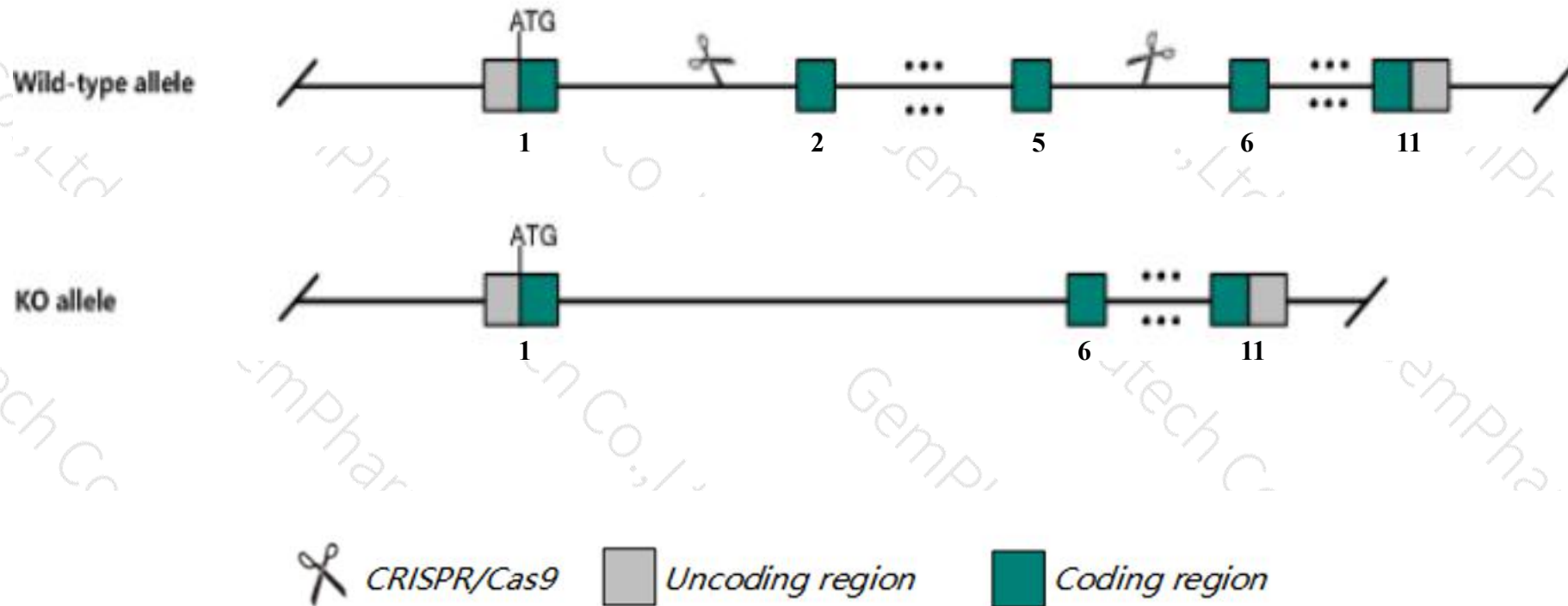
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Sohlh2* gene. The schematic diagram is as follows:



- The *Sohlh2* gene has 1 transcript. According to the structure of *Sohlh2* gene, exon2-exon5 of *Sohlh2*-201(ENSMUST00000029369.4) transcript is recommended as the knockout region. The region contains 479bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Sohlh2* 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, homozygous inactivation of this gene leads to female and male infertility due to defects in early oocyte and spermatogonial differentiation.
- The *Sohlh2* gene is located on the Chr3. 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)

## Sohlh2 spermatogenesis and oogenesis specific basic helix-loop-helix 2 [ *Mus musculus* (house mouse) ]

Gene ID: 74434, updated on 26-Jun-2020

### Summary



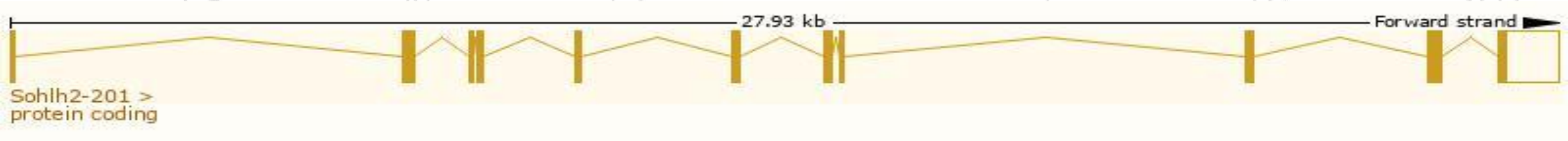
Official Symbol	Sohlh2 provided by <a href="#">MGI</a>
Official Full Name	spermatogenesis and oogenesis specific basic helix-loop-helix 2 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:1921684</a>
See related	<a href="#">Ensembl:ENSMUSG00000027794</a>
Gene type	protein coding
RefSeq 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	Sosf2; AI413378; 4933406N12Rik
Expression	Biased expression in testis adult (RPKM 4.3), placenta adult (RPKM 0.7) and 1 other tissue <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

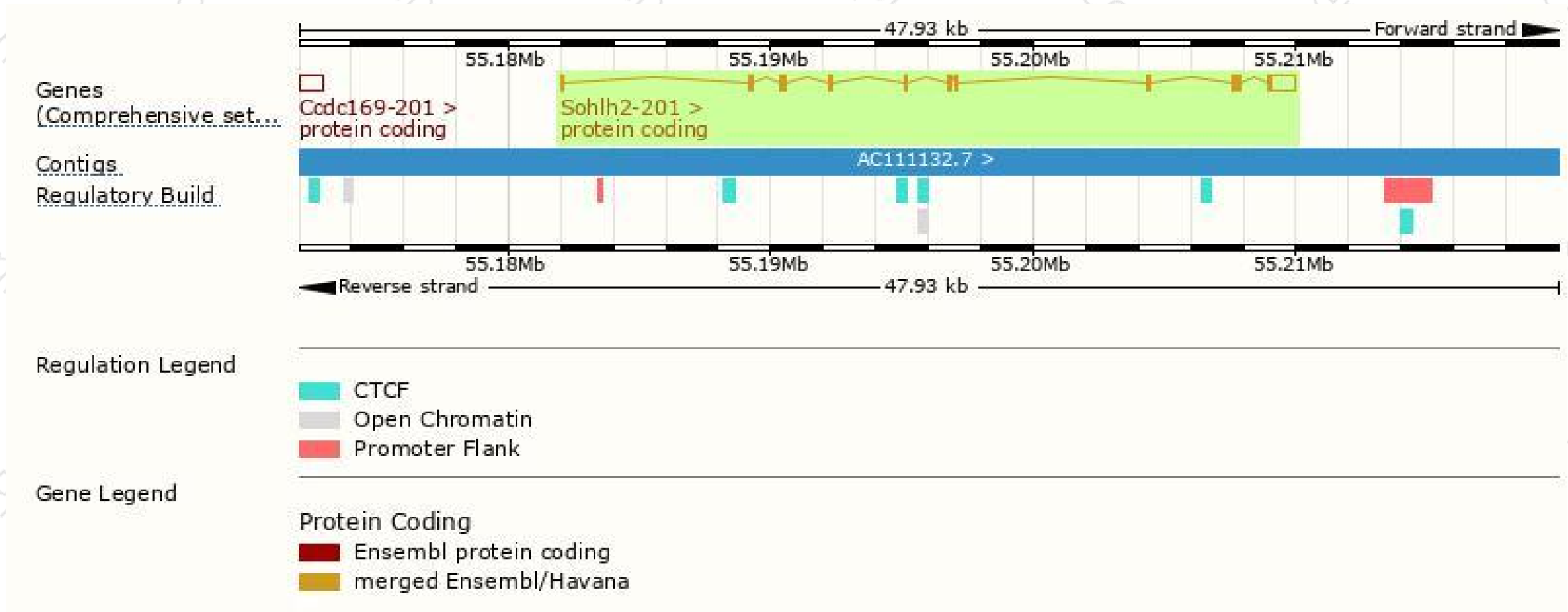
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Sohlh2-201	<a href="#">ENSMUST00000029369.4</a>	2364	<a href="#">467aa</a>	Protein coding	<a href="#">CCDS17358</a>	<a href="#">Q9D489</a>	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Sohlh2-201* transcript, the transcription is shown below:

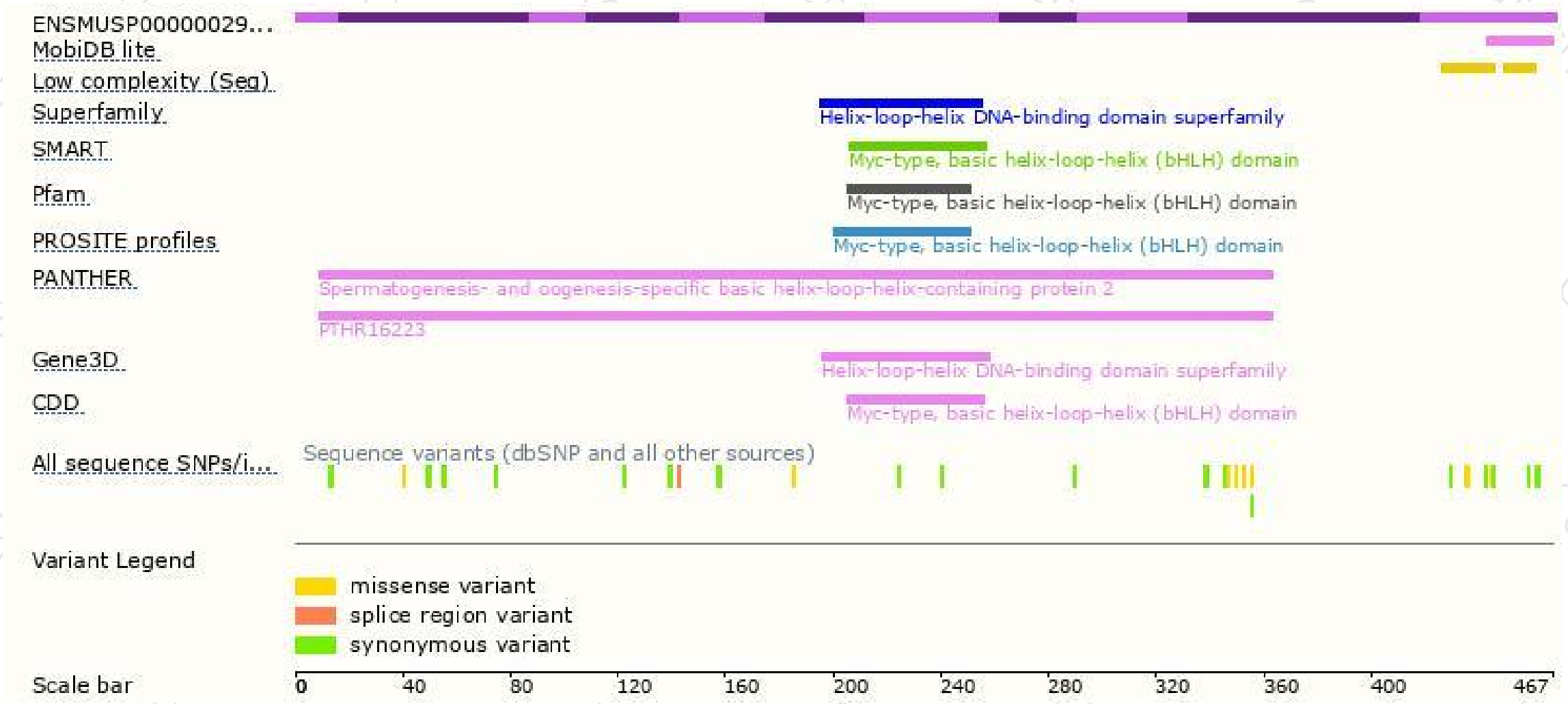


# Genomic location distribution

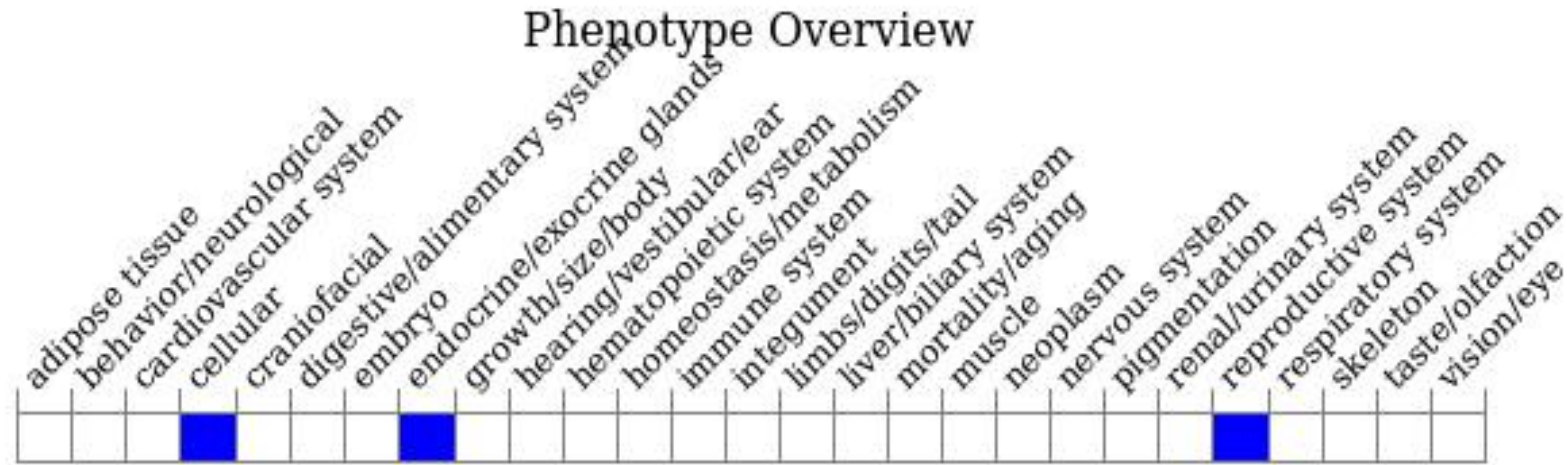




# Protein domain



# 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 inactivation of this gene leads to female and male infertility due to defects in early oocyte and spermatogonial differentiation.

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

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