

# *Nectin3* Cas9-KO Strategy

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**Reviewer: Shuang Zhang**

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# Project Overview

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**Project Name**

*Nectin3*

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**Project type**

**Cas9-KO**

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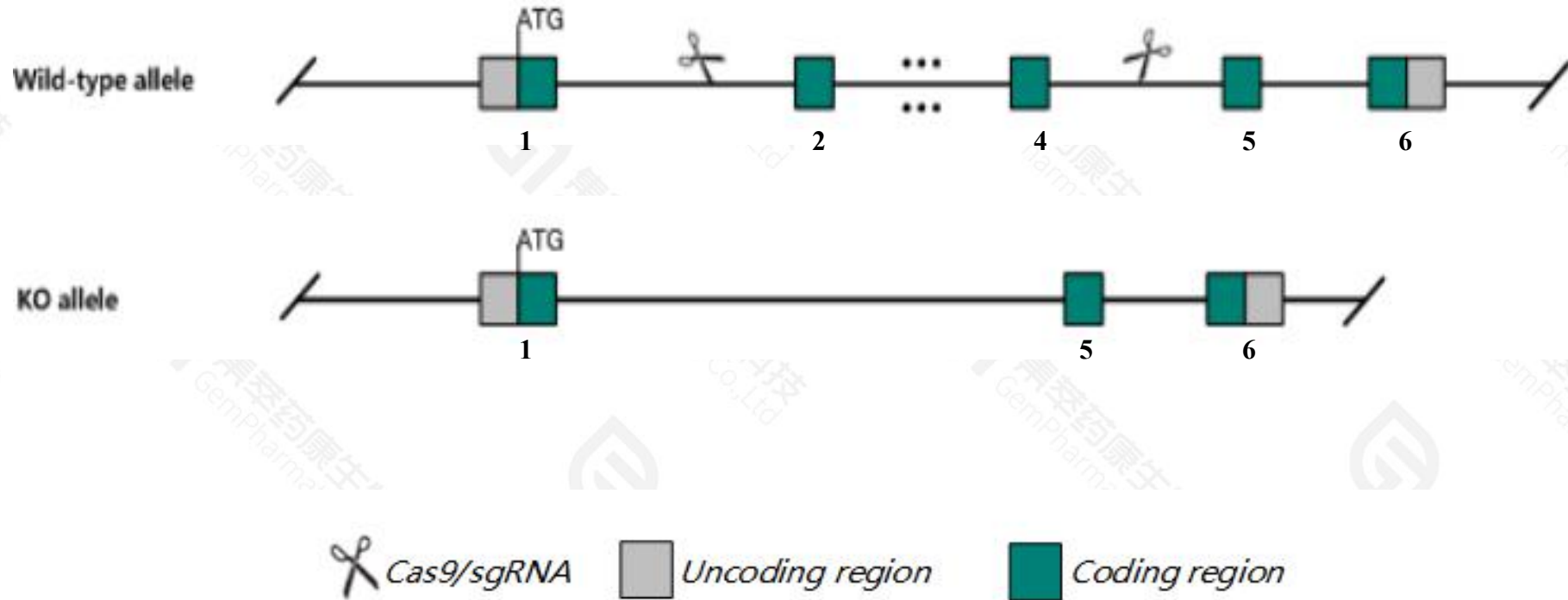
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Nectin3* gene. The schematic diagram is as follows:



- The *Nectin3* gene has 10 transcripts. According to the structure of *Nectin3* gene, exon2-exon4 of *Nectin3*-201(ENSMUST00000023334.15) transcript is recommended as the knockout region. The region contains 757bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Nectin3* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA 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 null mice exhibit male infertility and eye abnormalities including microphthalmia, absent vitreous body, abnormal ciliary body, retinal layers, and lenses.
- The *Nectin3* gene is located on the Chr16. 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.
- Transcript 204.205.206. 207.208 will not be destroyed.
- 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.

## Nectin3 nectin cell adhesion molecule 3 [Mus musculus (house mouse)]

Gene ID: 58998, updated on 21-Feb-2021

### Summary



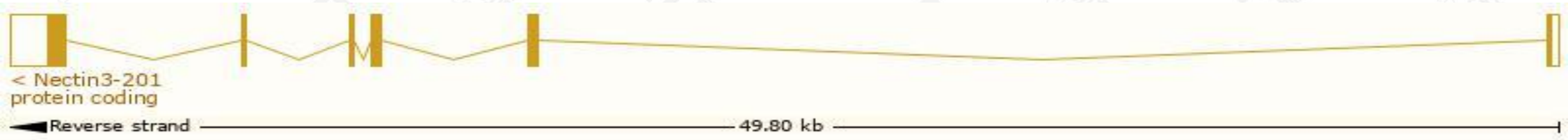
<b>Official Symbol</b>	Nectin3 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	nectin cell adhesion molecule 3 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1930171</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000022656</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	2610301B19Rik, 3000002N23Rik, 4921513D19Rik, AA407785, AU016832, AW538082, CD113, P, Pvr13, necti
<b>Expression</b>	Broad expression in testis adult (RPKM 13.1), CNS E11.5 (RPKM 4.7) and 19 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

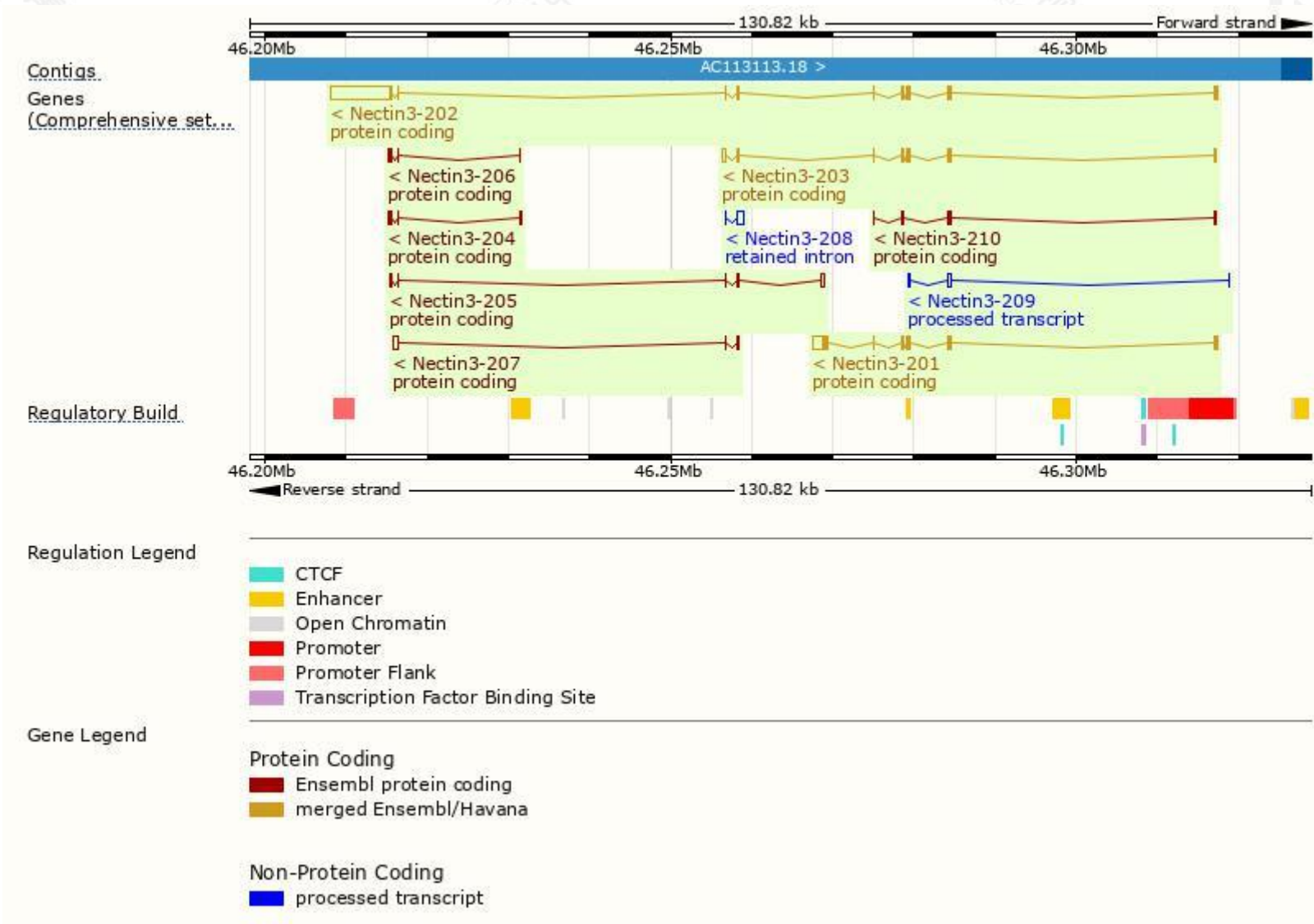
The gene has 10 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nectin3-202	<a href="#">ENSMUST00000023335.13</a>	9116	<a href="#">510aa</a>	Protein coding	<a href="#">CCDS28204</a>		TSL:1 , GENCODE basic , APPRIS ALT1 ,
Nectin3-201	<a href="#">ENSMUST00000023334.15</a>	3062	<a href="#">549aa</a>	Protein coding	<a href="#">CCDS28206</a>		TSL:1 , GENCODE basic , APPRIS P4 ,
Nectin3-203	<a href="#">ENSMUST00000096052.9</a>	1746	<a href="#">438aa</a>	Protein coding	<a href="#">CCDS28205</a>		TSL:1 , GENCODE basic ,
Nectin3-205	<a href="#">ENSMUST00000121245.8</a>	953	<a href="#">185aa</a>	Protein coding	-		TSL:3 , GENCODE basic ,
Nectin3-207	<a href="#">ENSMUST00000124602.2</a>	911	<a href="#">122aa</a>	Protein coding	-		CDS 5' incomplete , TSL:3 ,
Nectin3-210	<a href="#">ENSMUST00000149901.2</a>	732	<a href="#">244aa</a>	Protein coding	-		CDS 5' and 3' incomplete , TSL:3 ,
Nectin3-206	<a href="#">ENSMUST00000121803.8</a>	541	<a href="#">100aa</a>	Protein coding	-		TSL:2 , GENCODE basic ,
Nectin3-204	<a href="#">ENSMUST00000119941.2</a>	529	<a href="#">93aa</a>	Protein coding	-		TSL:3 , GENCODE basic ,
Nectin3-209	<a href="#">ENSMUST00000133935.2</a>	737	No protein	Processed transcript	-		TSL:3 ,
Nectin3-208	<a href="#">ENSMUST00000132089.2</a>	752	No protein	Retained intron	-		TSL:2 ,

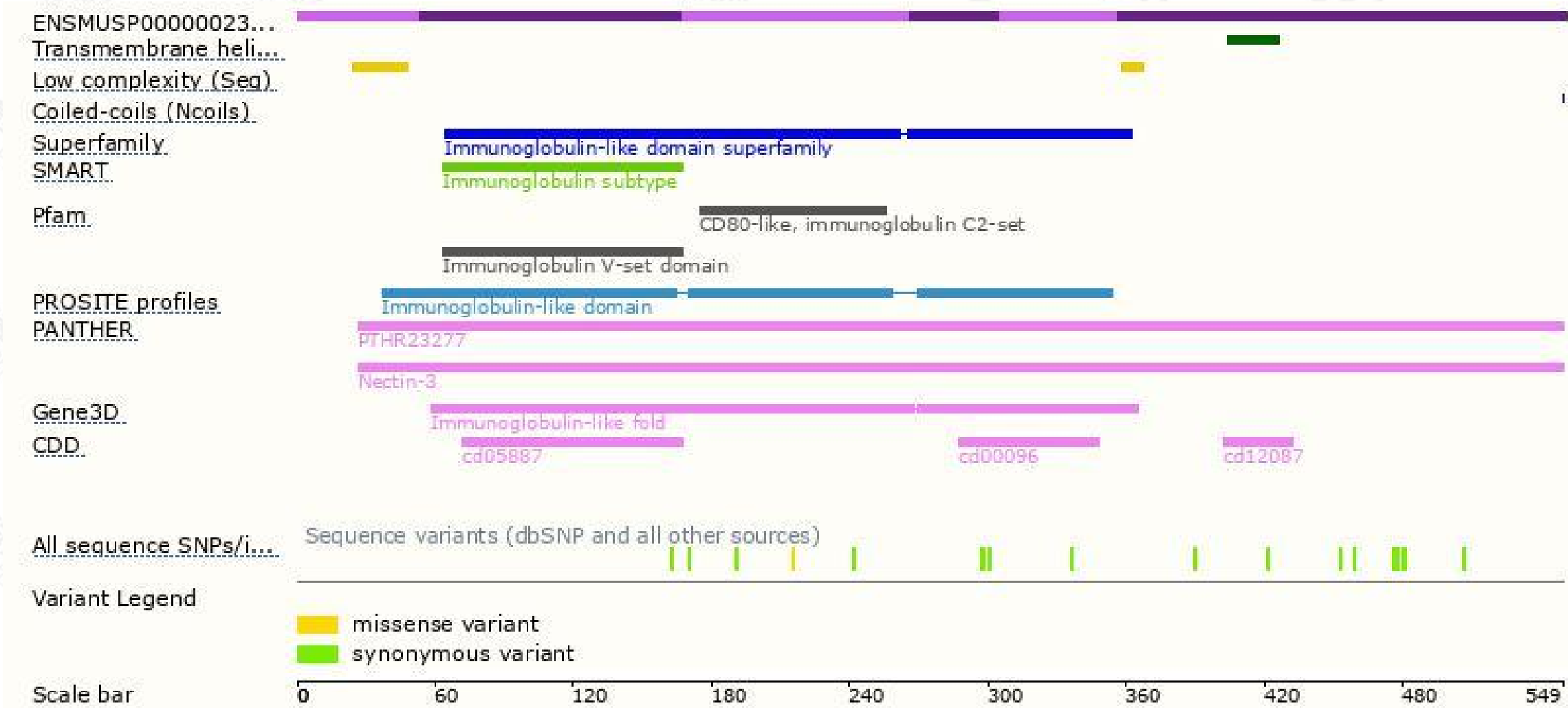
The strategy is based on the design of *Nectin3-201* transcript,the transcription is shown below:



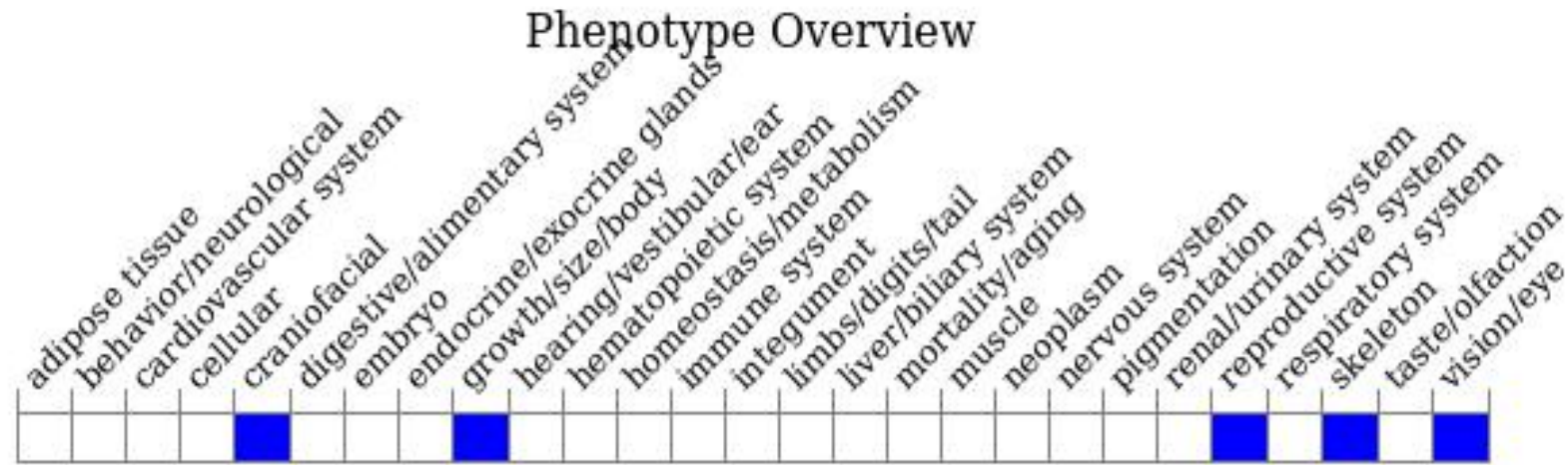
# Genomic location distribution



# 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 null mice exhibit male infertility and eye abnormalities including microphthalmia, absent vitreous body, abnormal ciliary body, retinal layers, and lenses.

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

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