

# *Anks3* Cas9-KO Strategy

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**Reviewer:**

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**Design Date:**

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# Project Overview

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**Project Name**

*Anks3*

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**Project type**

**Cas9-KO**

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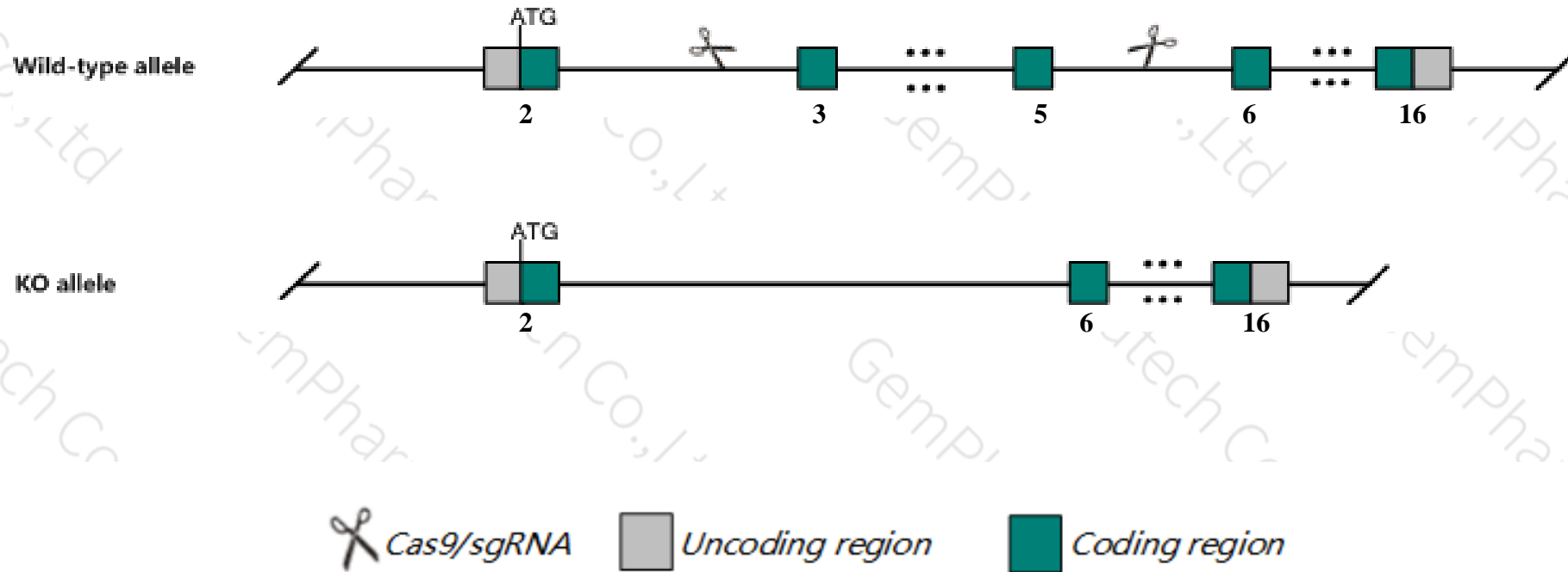
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Anks3* gene. The schematic diagram is as follows:



- The *Anks3* gene has 13 transcripts. According to the structure of *Anks3* gene, exon3-exon5 of *Anks3-201* (ENSMUST00000023157.5) transcript is recommended as the knockout region. The region contains 403bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Anks3* 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.

- Transcript *Anks3*-203/204/205/207/208/209/210/213 may not be affected.
- The *Anks3* 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.
- 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.

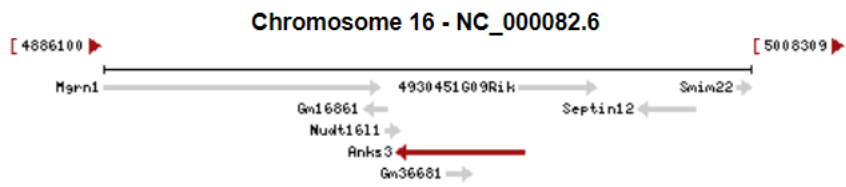
# Gene information (NCBI)

## Anks3 ankyrin repeat and sterile alpha motif domain containing 3 [ *Mus musculus* (house mouse) ]

Gene ID: 72615, updated on 28-Oct-2019

### Summary

- Official Symbol** Anks3 provided by [MGI](#)
- Official Full Name** ankyrin repeat and sterile alpha motif domain containing 3 provided by [MGI](#)
- Primary source** [MGI:MGI:1919865](#)
- See related** [Ensembl:ENSMUSG00000022515](#)
- 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** C81345; mKIAA1977; 2700067D09Rik
- Expression** Ubiquitous expression in CNS E14 (RPKM 12.6), CNS E18 (RPKM 12.3) and 28 other tissues [See more](#)
- Orthologs** [human](#) [all](#)



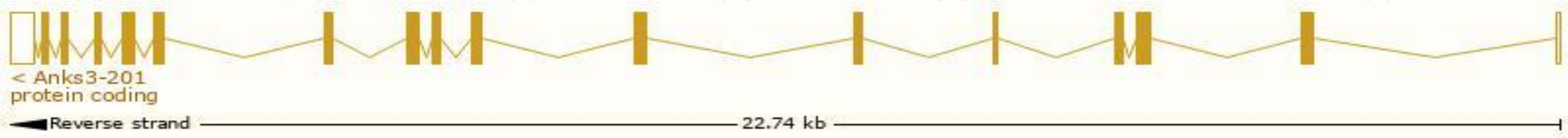


# Transcript information (Ensembl)

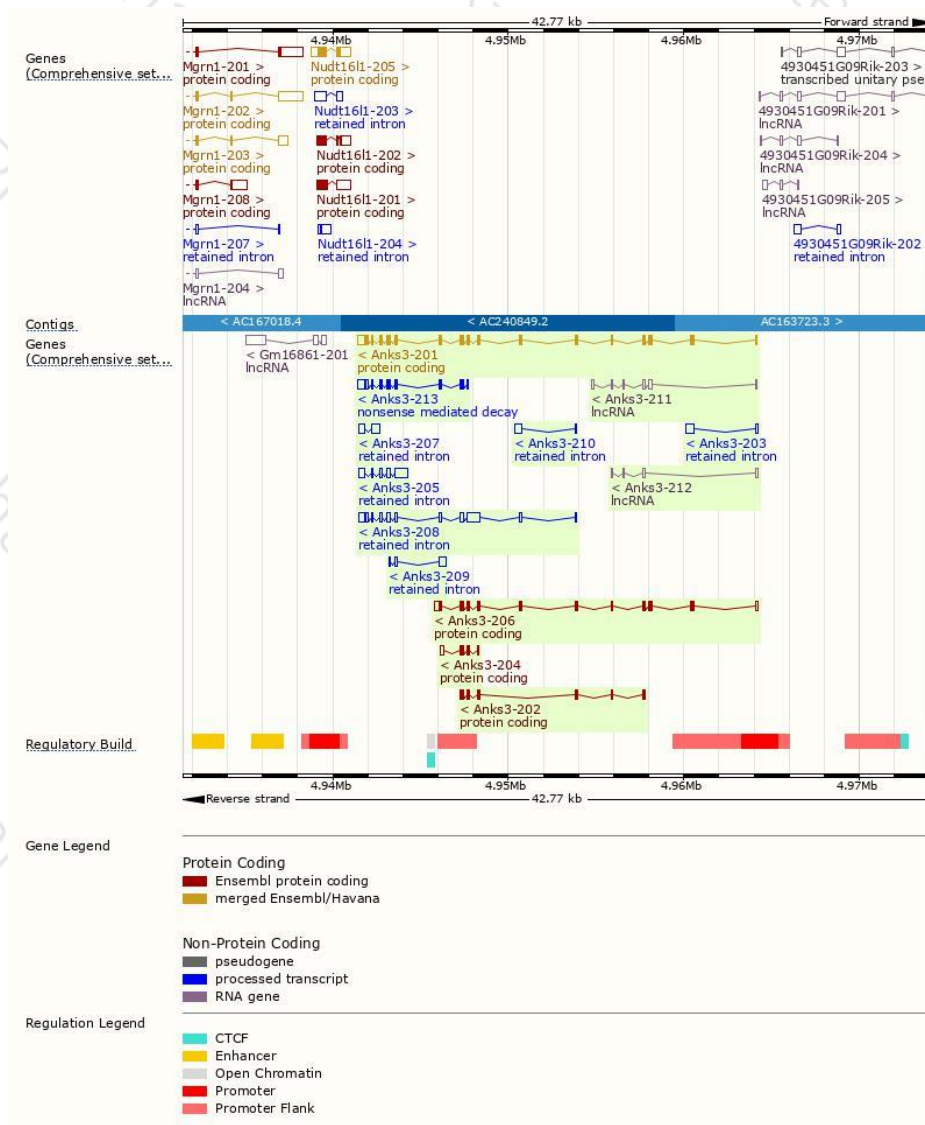
The gene has 13 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Translation ID	Biotype	CCDS	UniProt	Flags
Anks3-201	<a href="#">ENSMUST00000023157.5</a>	2398	<a href="#">655aa</a>	<a href="#">ENSMUSP00000023157.4</a>	Protein coding	<a href="#">CCDS27928</a>	<a href="#">Q9CZK6</a>	TSL:1 GENCODE basic APPRIS P1
Anks3-206	<a href="#">ENSMUST000000229765.1</a>	1779	<a href="#">469aa</a>	<a href="#">ENSMUSP000000155302.1</a>	Protein coding	-	<a href="#">Q80XC9</a>	GENCODE basic
Anks3-202	<a href="#">ENSMUST000000229017.1</a>	739	<a href="#">246aa</a>	<a href="#">ENSMUSP000000155720.1</a>	Protein coding	-	<a href="#">A0A2R8VI44</a>	CDS 5' and 3' incomplete
Anks3-204	<a href="#">ENSMUST000000229272.1</a>	608	<a href="#">140aa</a>	<a href="#">ENSMUSP000000155421.1</a>	Protein coding	-	<a href="#">A0A2R8VKE4</a>	CDS 5' incomplete
Anks3-213	<a href="#">ENSMUST000000231036.1</a>	1289	<a href="#">266aa</a>	<a href="#">ENSMUSP000000155247.1</a>	Nonsense mediated decay	-	<a href="#">A0A2R8VHN6</a>	CDS 5' incomplete
Anks3-208	<a href="#">ENSMUST000000230083.1</a>	2144	No protein	-	Retained intron	-	-	-
Anks3-205	<a href="#">ENSMUST000000229477.1</a>	1372	No protein	-	Retained intron	-	-	-
Anks3-207	<a href="#">ENSMUST000000229898.1</a>	799	No protein	-	Retained intron	-	-	-
Anks3-209	<a href="#">ENSMUST000000230466.1</a>	592	No protein	-	Retained intron	-	-	-
Anks3-203	<a href="#">ENSMUST000000229077.1</a>	537	No protein	-	Retained intron	-	-	-
Anks3-210	<a href="#">ENSMUST000000230493.1</a>	476	No protein	-	Retained intron	-	-	-
Anks3-211	<a href="#">ENSMUST000000230721.1</a>	668	No protein	-	lncRNA	-	-	-
Anks3-212	<a href="#">ENSMUST000000230771.1</a>	360	No protein	-	lncRNA	-	-	-

The strategy is based on the design of *Anks3-201* transcript,The transcription is shown below

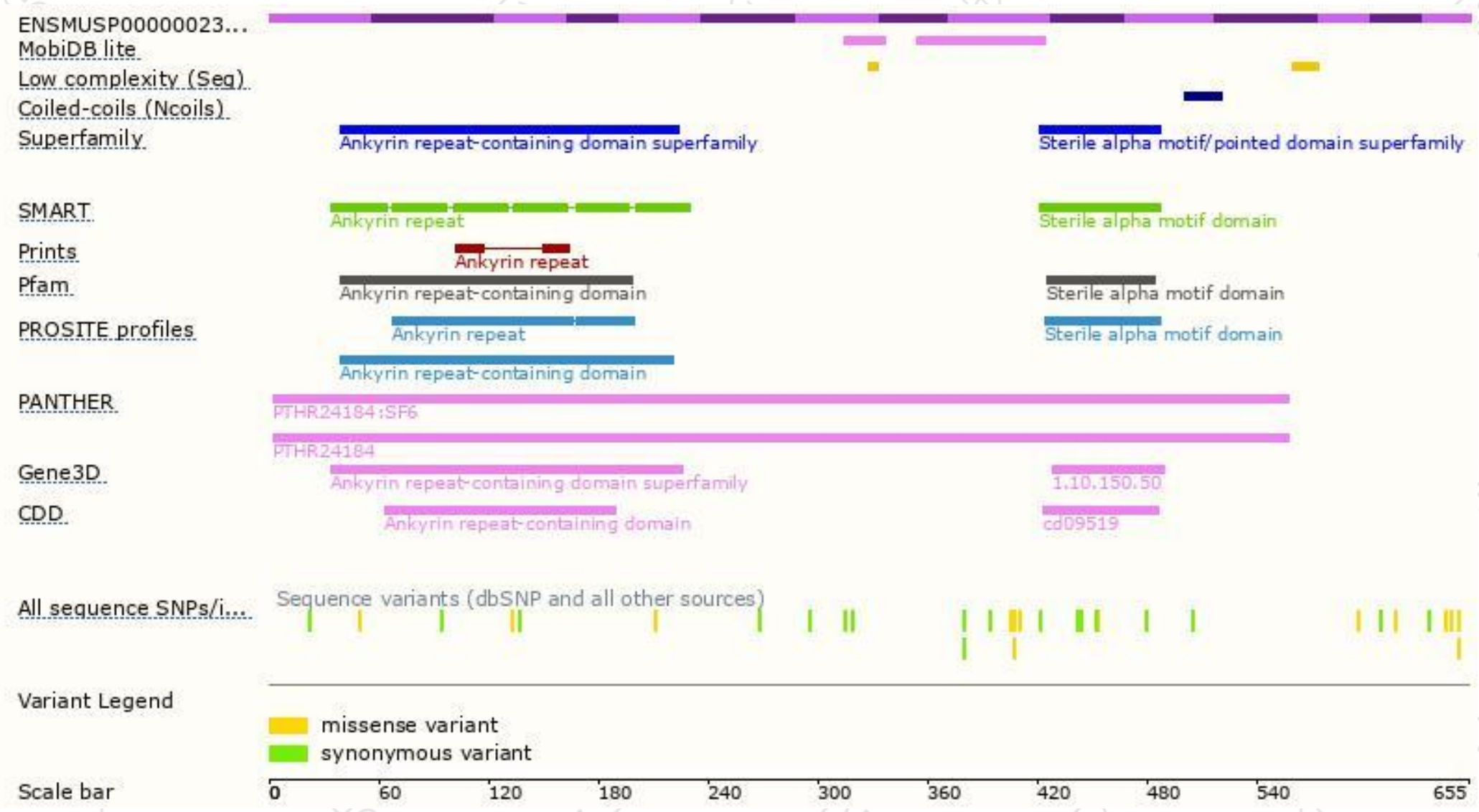


# Genomic location distribution



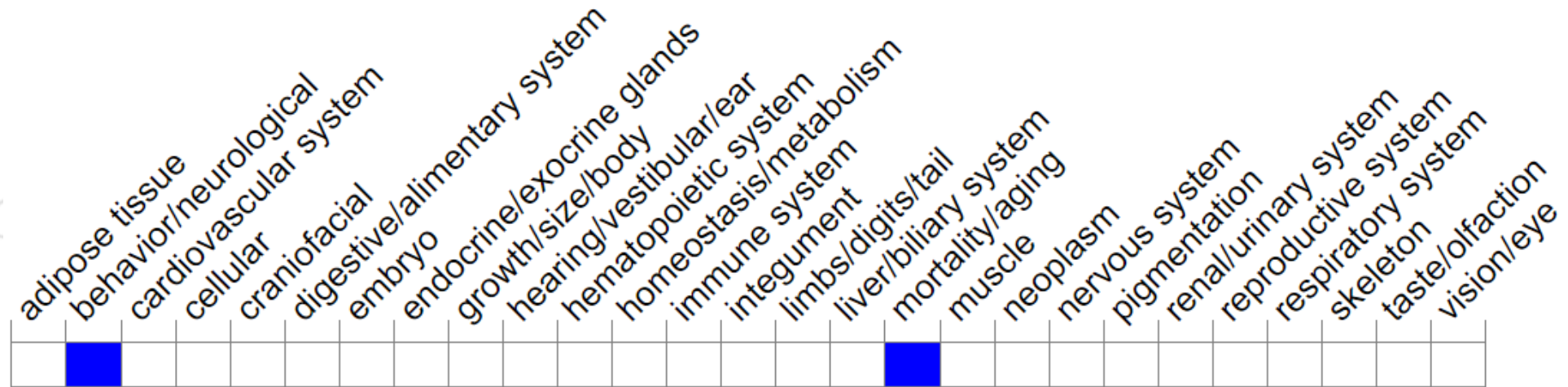


# Protein domain



# Mouse phenotype description(MGI)

## Phenotype Overview ?



*Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).*

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

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