

# ***Hsd17b4 Cas9-KO Strategy***

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**Reviewer:**

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# Project Overview

**Project Name**

*Hsd17b4*

**Project type**

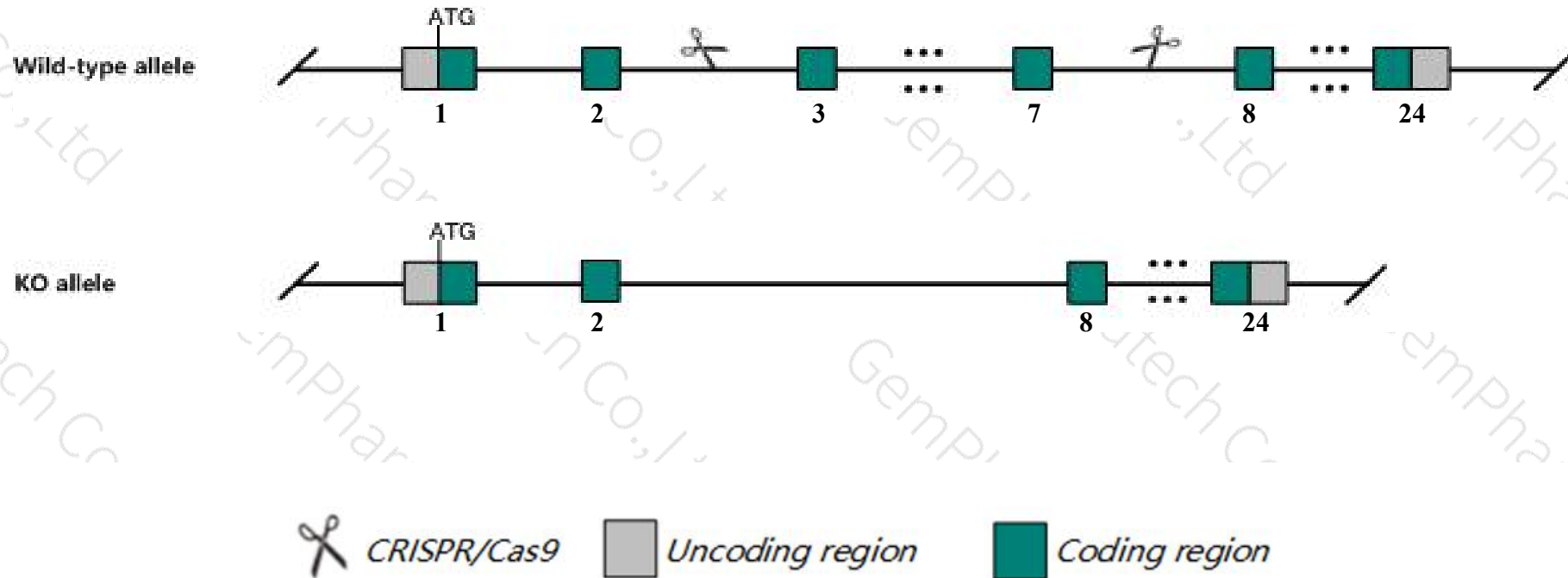
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Hsd17b4* gene. The schematic diagram is as follows:



- The *Hsd17b4* gene has 4 transcripts. According to the structure of *Hsd17b4* gene, exon3-exon7 of *Hsd17b4-201* (ENSMUST00000025385.6) transcript is recommended as the knockout region. The region contains 322bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Hsd17b4* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for disruptions in this gene have abnormalities in fatty acid metabolism, retarded growth, abnormal bile salt composition, impaired coordination, demyelination and premature death.
- The *Hsd17b4* 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.



# Gene information (NCBI)

## Hsd17b4 hydroxysteroid (17-beta) dehydrogenase 4 [Mus musculus (house mouse)]

Gene ID: 15488, updated on 7-Apr-2019

### Summary



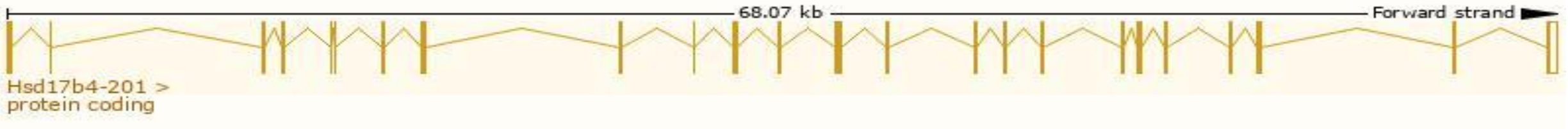
<b>Official Symbol</b>	Hsd17b4 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	hydroxysteroid (17-beta) dehydrogenase 4 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:105089</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000024507</a>
<b>Gene type</b>	protein 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>	17-beta-HSD, 17[b]-HSD, DBP, MFE-2, MFP2, MPF-2, Mfp-2, perMFE-2
<b>Expression</b>	Ubiquitous expression in liver adult (RPKM 28.9), placenta adult (RPKM 27.7) and 28 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

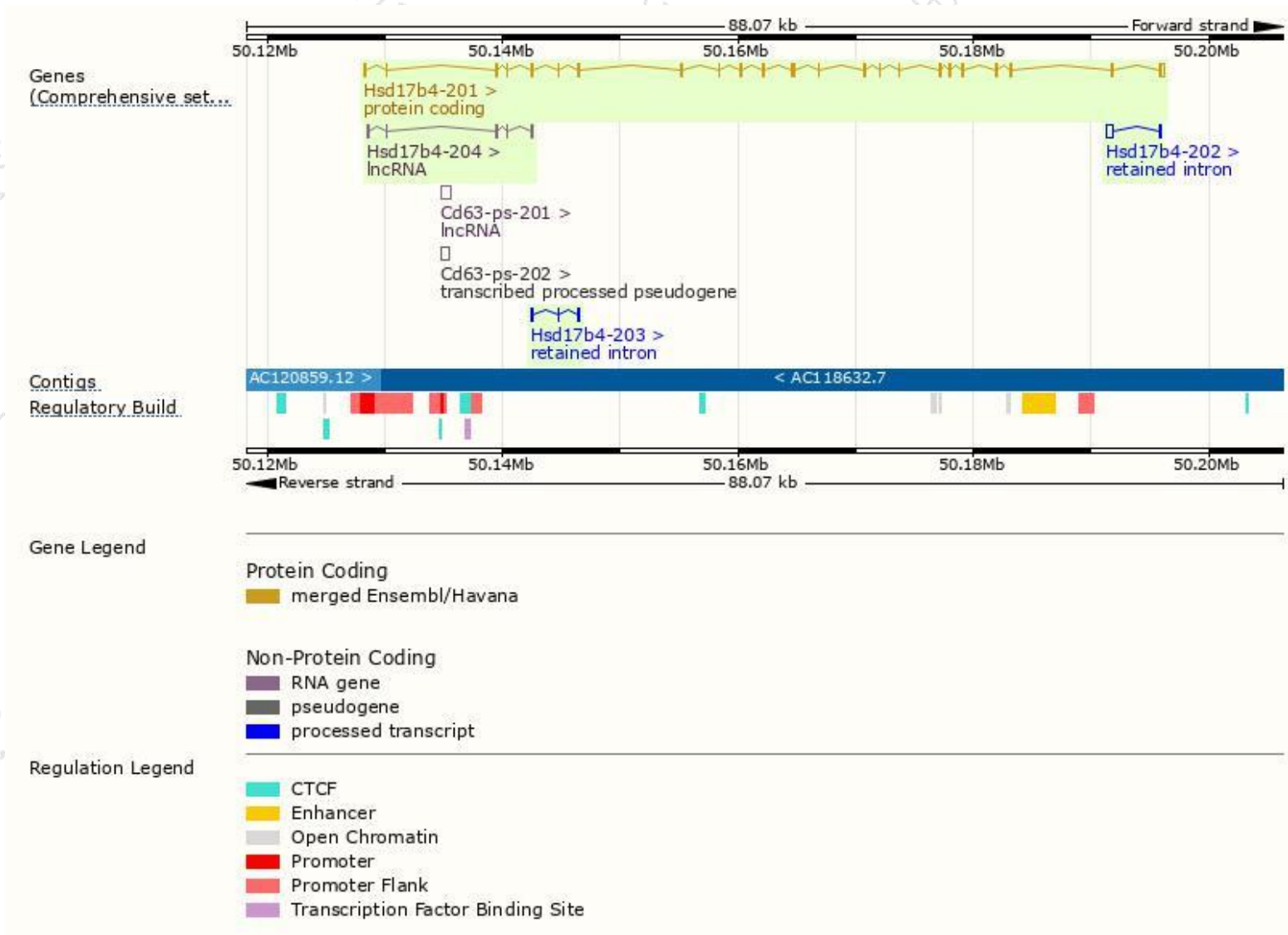
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hsd17b4-201	<a href="#">ENSMUST00000025385.6</a>	2684	<a href="#">735aa</a>	Protein coding	<a href="#">CCDS29242</a>	<a href="#">P51660</a>	TSL:1 GENCODE basic APPRIS P1
Hsd17b4-202	<a href="#">ENSMUST00000127325.1</a>	634	No protein	Retained intron	-	-	TSL:1
Hsd17b4-203	<a href="#">ENSMUST00000139502.1</a>	536	No protein	Retained intron	-	-	TSL:2
Hsd17b4-204	<a href="#">ENSMUST00000152388.1</a>	394	No protein	lncRNA	-	-	TSL:3

The strategy is based on the design of *Hsd17b4-201* transcript,The transcription is shown below



# Genomic location distribution

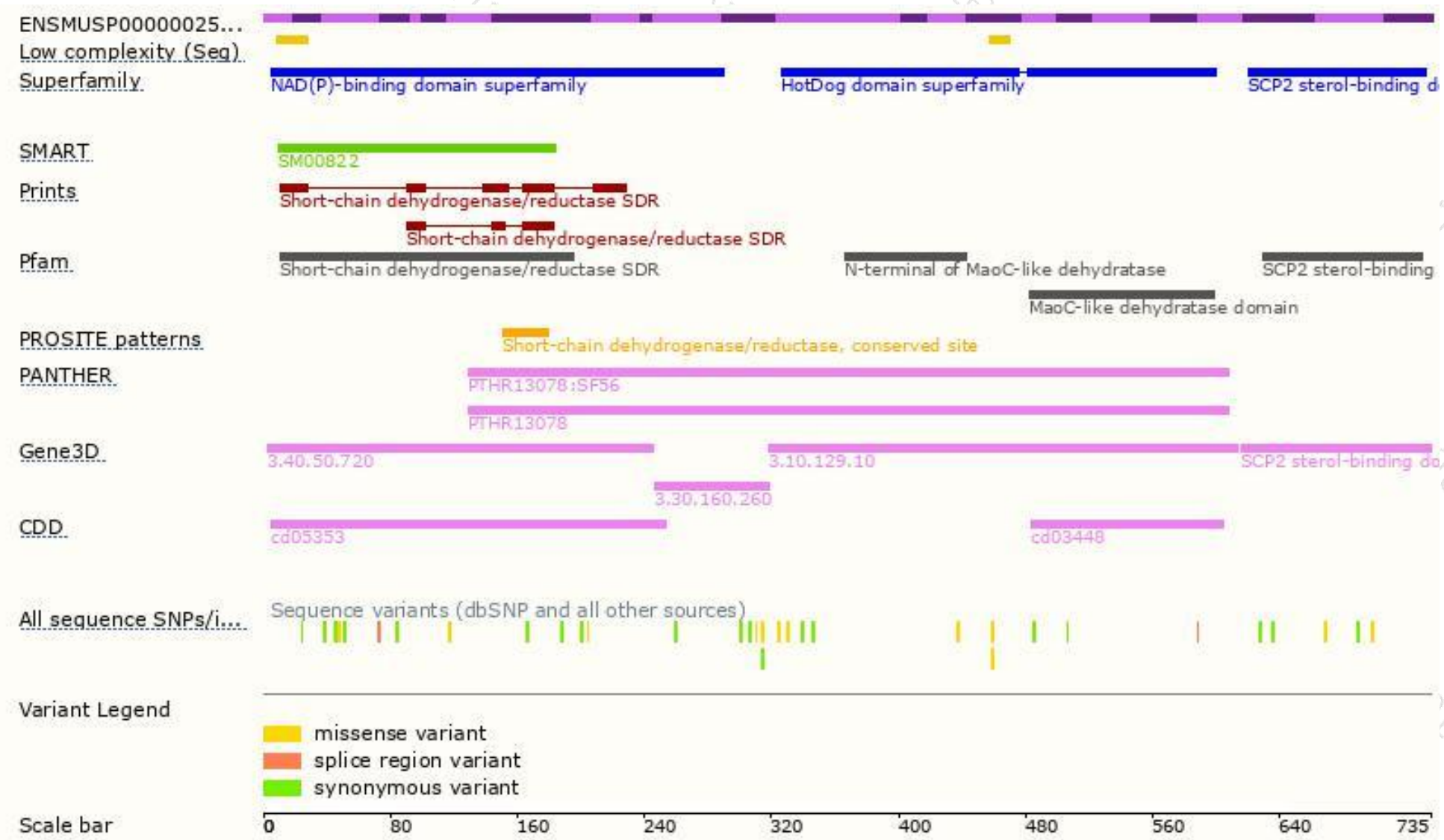




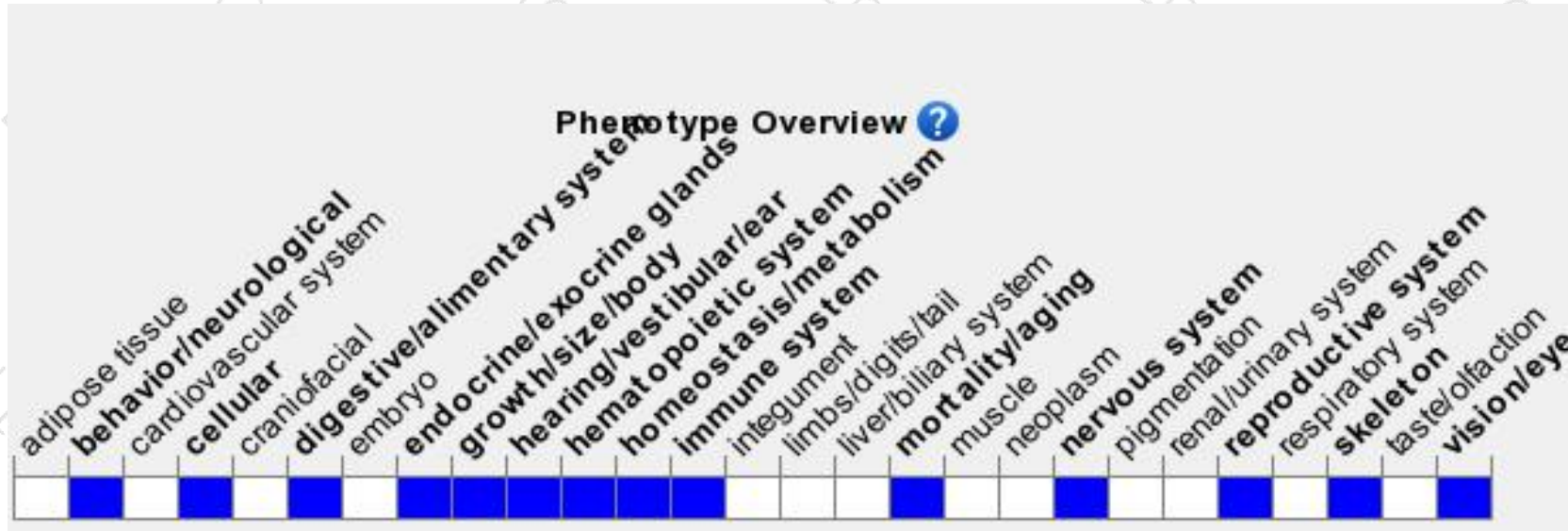
# Protein domain



集萃药康  
GemPharmatech



# 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 disruptions in this gene have abnormalities in fatty acid metabolism, retarded growth, abnormal bile salt composition, impaired coordination, demyelination and premature death.

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

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