



Pdhal Cas9-CKO Strategy

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Reviewer: Shilei Zhu

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Project Overview

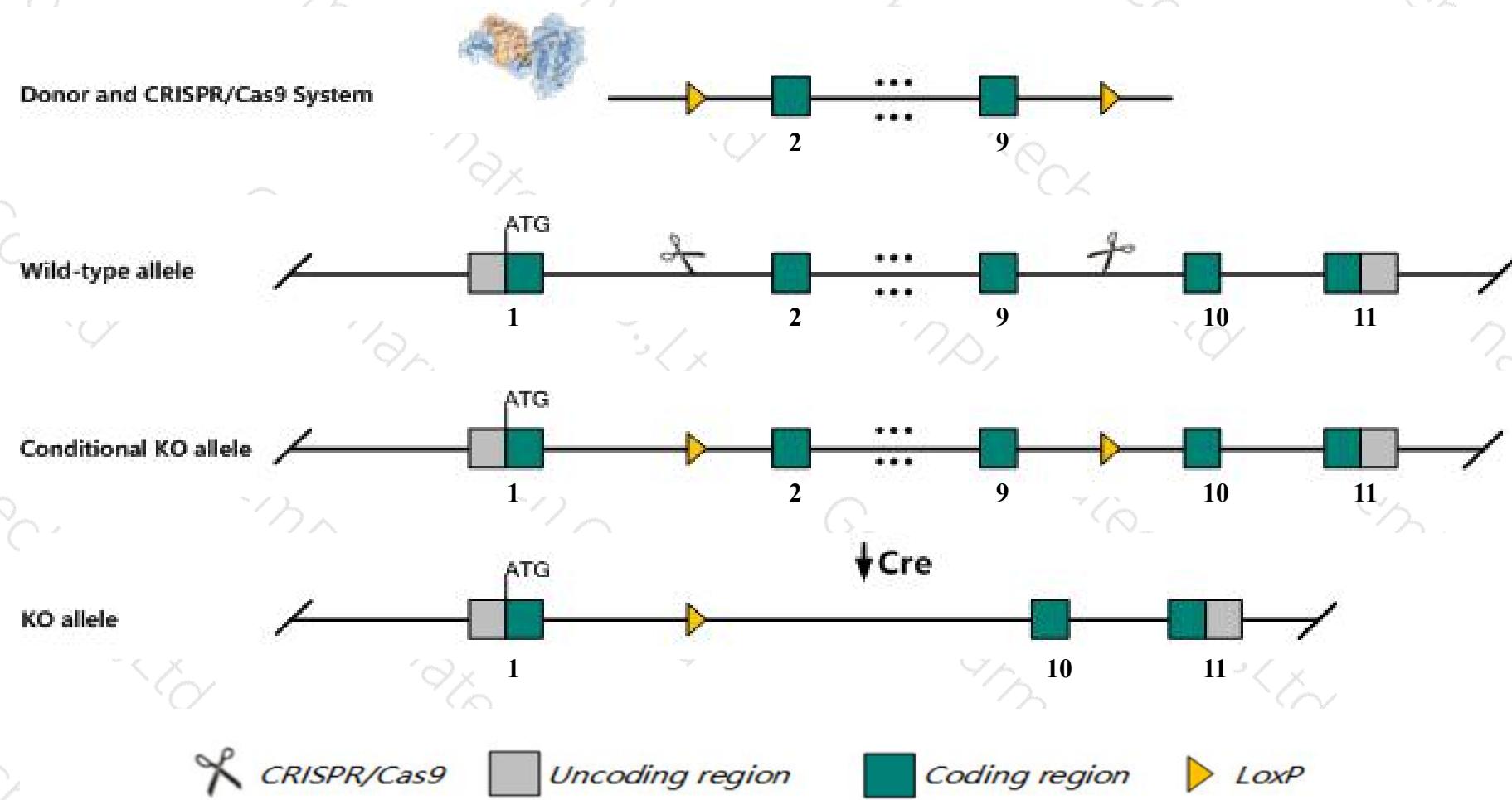
Project Name**Pdhal**

Project type**Cas9-CKO**

Strain background**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Pdhα1* gene. The schematic diagram is as follows:



Technical routes

- The *Pdhal* gene has 4 transcripts. According to the structure of *Pdhal* gene, exon2-exon9 of *Pdhal*-201 (ENSMUST00000033662.8) transcript is recommended as the knockout region. The region contains 842bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Pdhal* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.



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Notice

- According to the existing MGI data, Mice homozygous for a knock-out allele die by E12.5.
- The *Pdha1* gene is located on the ChrX. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.



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Gene information (NCBI)

Pdha1 pyruvate dehydrogenase E1 alpha 1 [Mus musculus (house mouse)]

Gene ID: 18597, updated on 7-Apr-2019

Summary



Official Symbol Pdha1 provided by [MGI](#)

Official Full Name pyruvate dehydrogenase E1 alpha 1 provided by [MGI](#)

Primary source [MGI:MGI:97532](#)

See related [Ensembl:ENSMUSG00000031299](#)

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 Pdha-1

Expression Ubiquitous expression in heart adult (RPKM 147.9), kidney adult (RPKM 68.4) and 26 other tissues [See more](#)

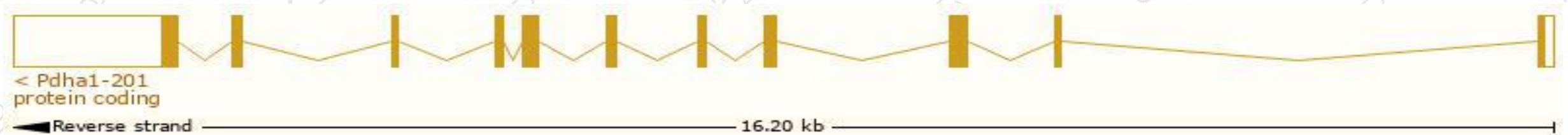
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

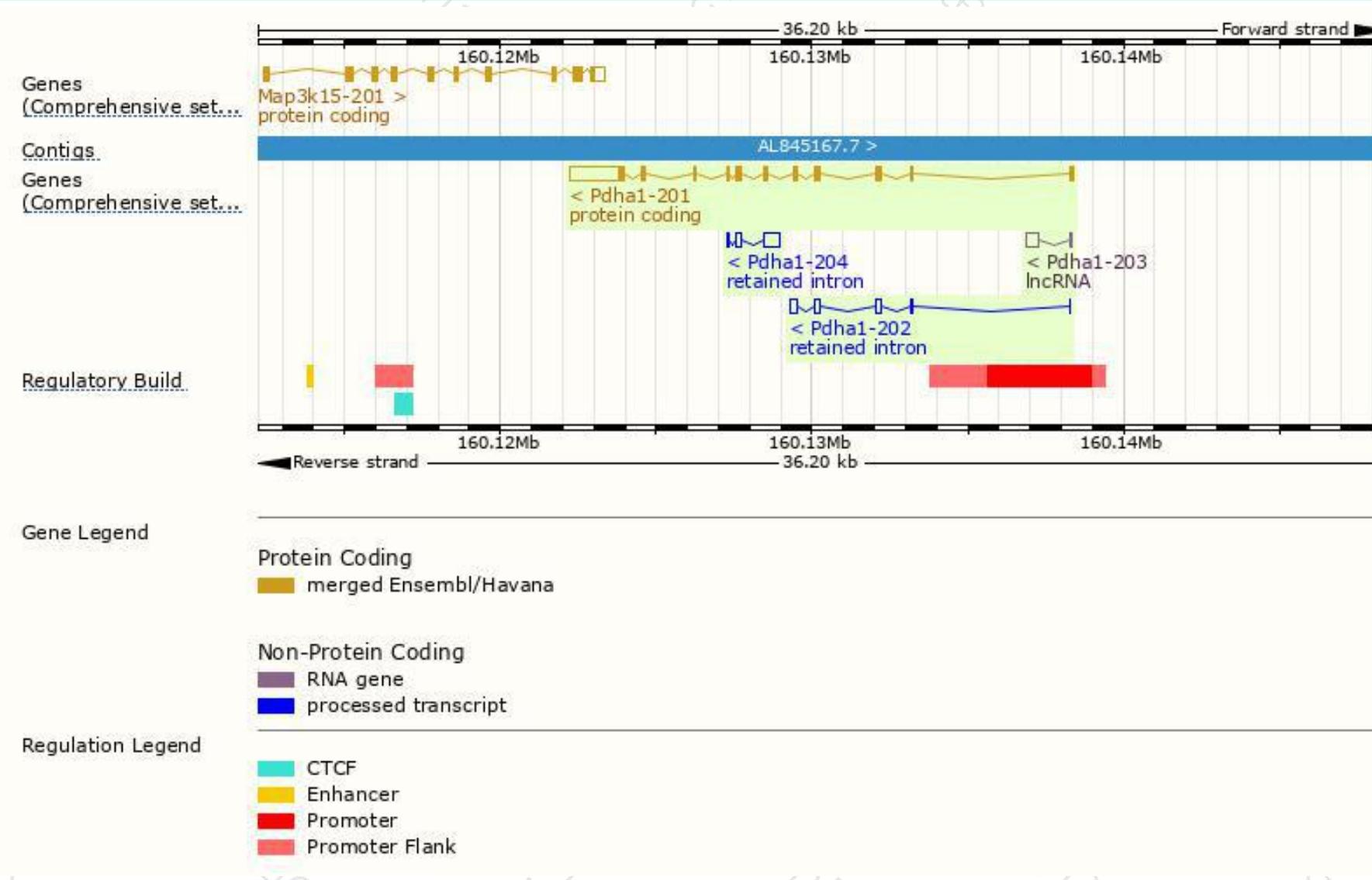
The gene has 4 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Pdha1-201	ENSMUST00000033662.8	2848	390aa	Protein coding	CCDS41195	P35486 Q3UFJ3	TSL:1 GENCODE basic APPRIS P1
Pdha1-204	ENSMUST00000156531.1	729	No protein	Retained intron	-	-	TSL:5
Pdha1-202	ENSMUST00000126440.1	629	No protein	Retained intron	-	-	TSL:2
Pdha1-203	ENSMUST00000127535.1	457	No protein	lncRNA	-	-	TSL:3

The strategy is based on the design of *Pdha1-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain

ENSMUSP000000033...

Low complexity (Seq)

TIGRFAM

Superfamily

Pfam

PANTHER

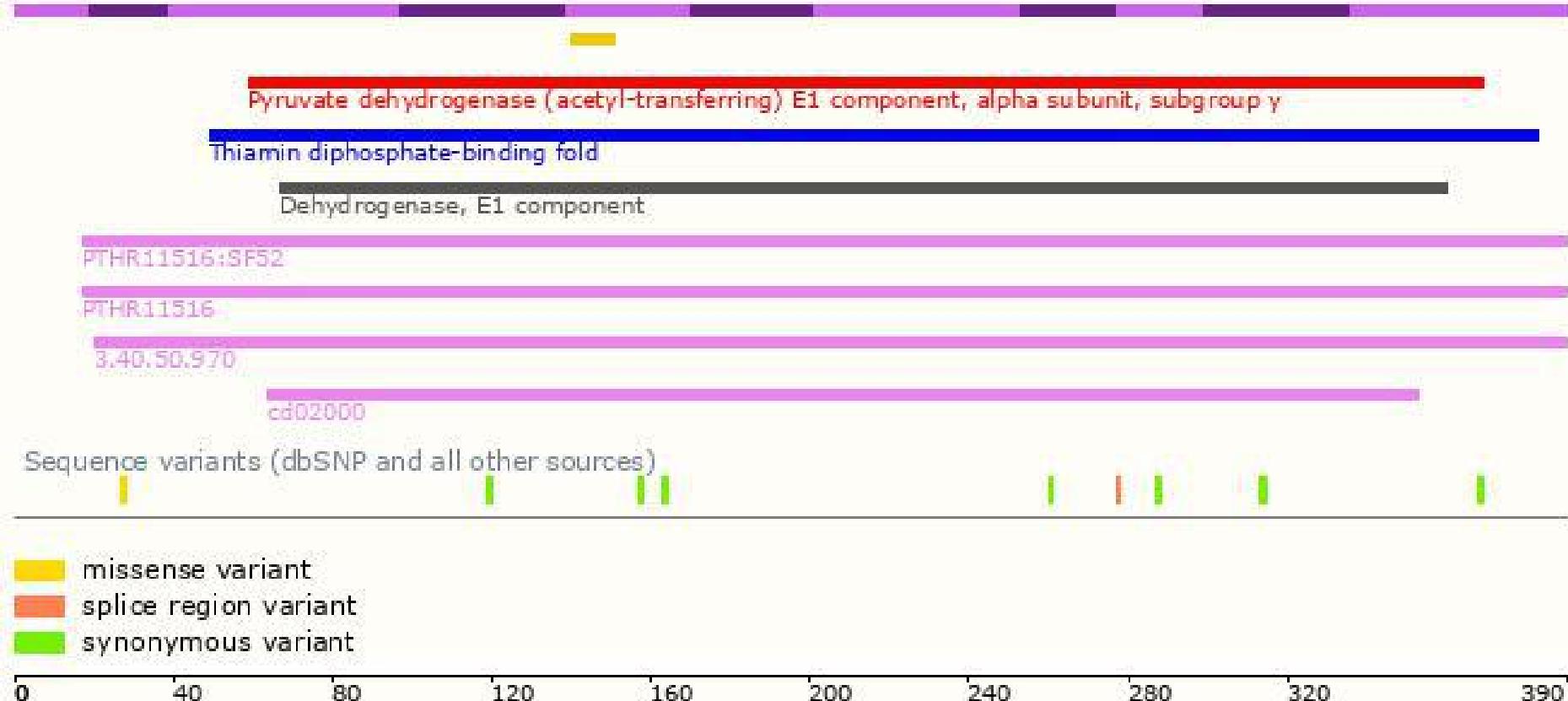
Gene3D

CDD

All sequence SNPs/i...

Variant Legend

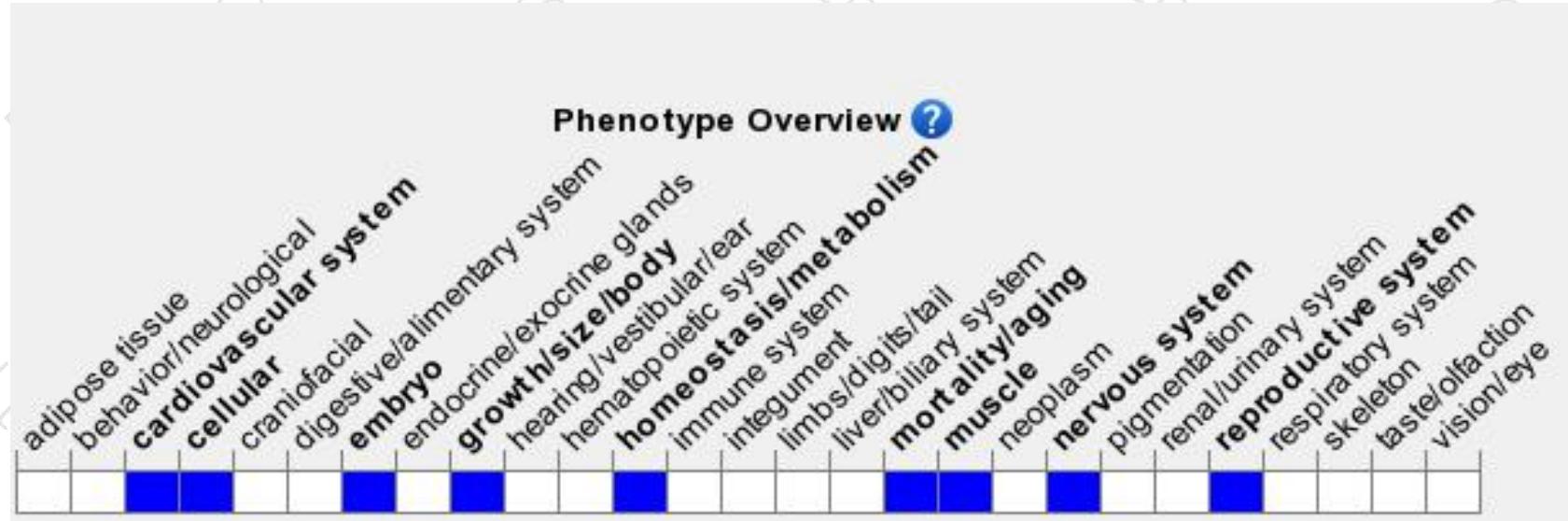
Scale bar





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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 die by E12.5.



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

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