

Pdha1 Cas9-KO Strategy

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Date: 2019/12/23

Project Overview

Project Name

Pdha1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Pdha1* gene has 4 transcripts. According to the structure of *Pdha1* gene, exon2-exon9 of *Pdha1-201* (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 *Pdha1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele die by E12.5.
- The *Pdhal* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

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

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

Summary



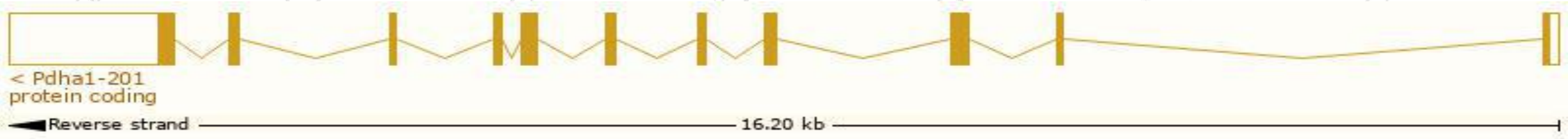
Official Symbol	Pdha1 provided by MGI
Official Full Name	pyruvate dehydrogenase E1 alpha 1 provided by MGI
Primary source	MGI:MGI:97532
See related	Ensembl:ENSMUSG00000031299
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; 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)

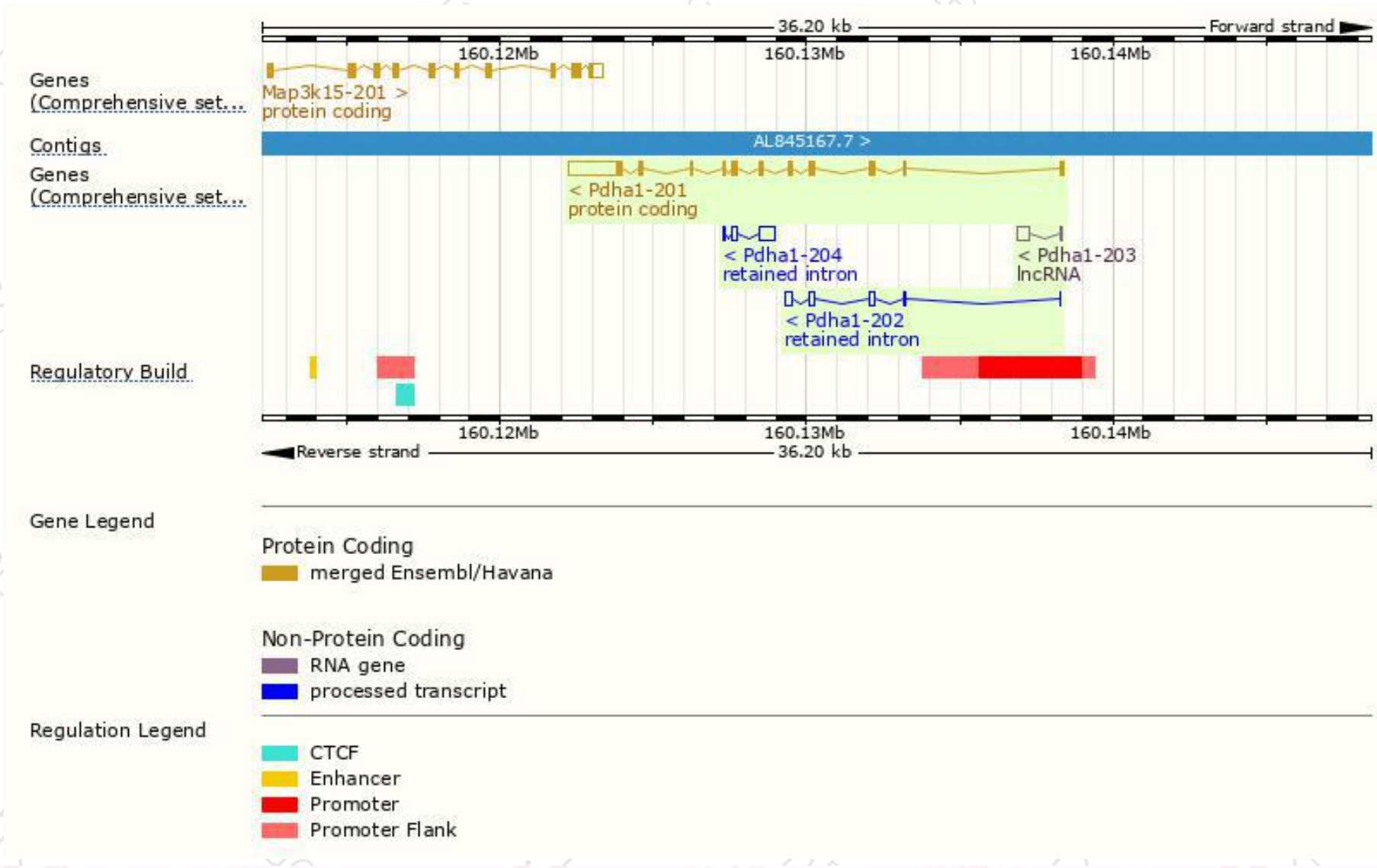
The gene has 4 transcripts,all transcripts are shown below:

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

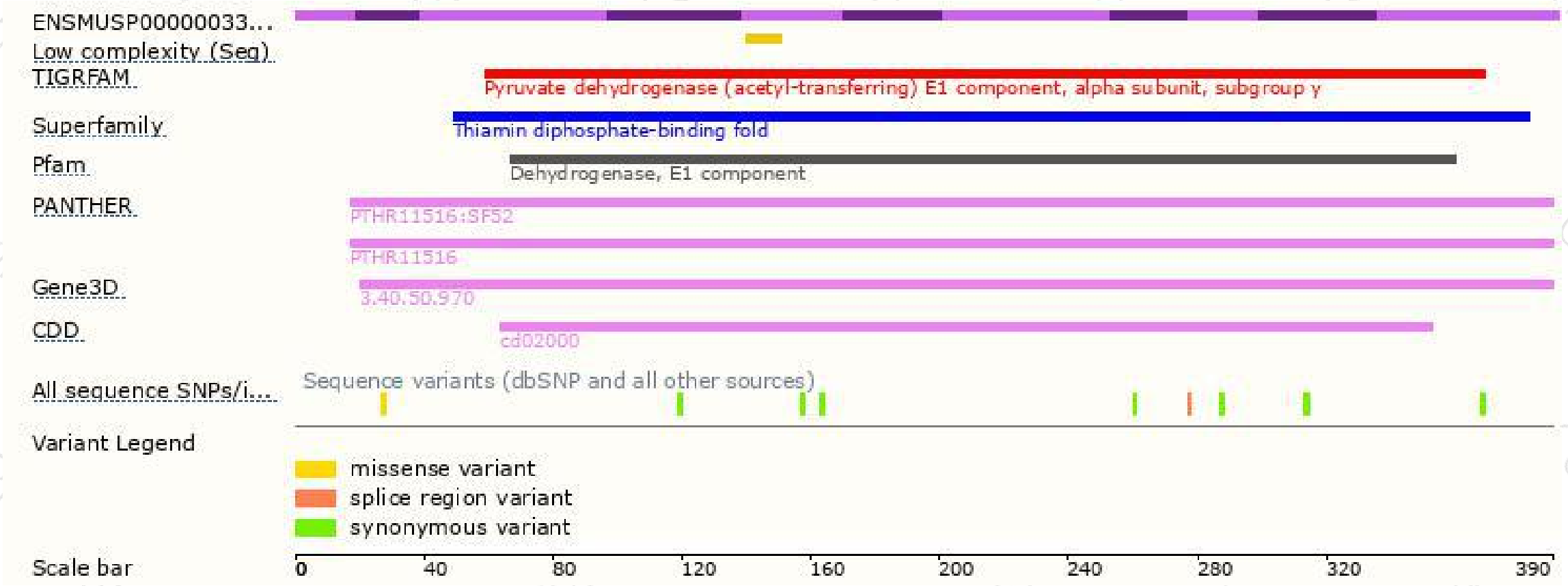
The strategy is based on the design of *Pdha1-201* transcript,The transcription is shown below



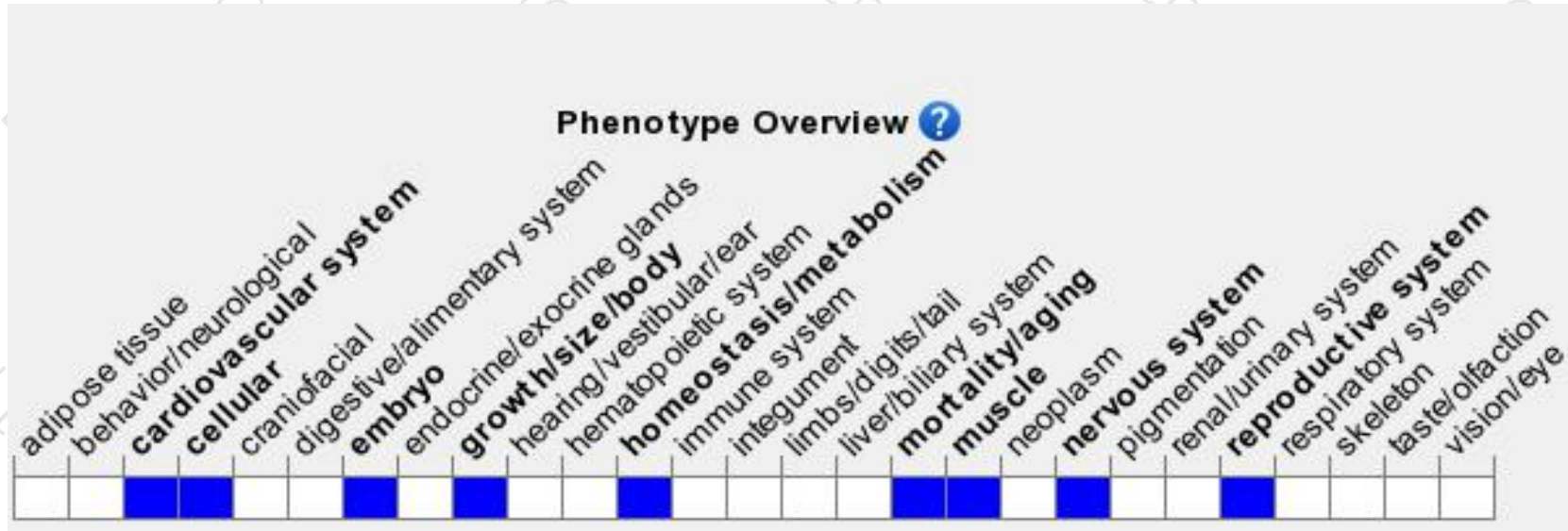
Genomic location distribution



Protein domain



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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