

***Enoph1* Cas9-KO Strategy**

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Reviewer:

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Design Date:

2020-2-27

Project Overview

Project Name

Enoph1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Enoph1* gene. The schematic diagram is as follows:



- The *Enoph1* gene has 4 transcripts. According to the structure of *Enoph1* gene, exon2-exon5 of *Enoph1*-202 (ENSMUST00000169390.7) transcript is recommended as the knockout region. The region contains 553bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Enoph1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Enoph1* gene is located on the Chr5. 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)

Enoph1 enolase-phosphatase 1 [Mus musculus (house mouse)]

Gene ID: 67870, updated on 31-Jan-2019

Summary



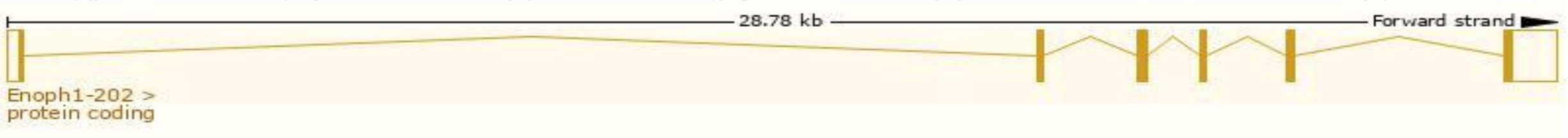
Official Symbol	Enoph1 provided by MGI
Official Full Name	enolase-phosphatase 1 provided by MGI
Primary source	MGI:MGI:1915120
See related	Ensembl:ENSMUSG00000029326
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2310057D15Rik, BB183658, C81437
Expression	Ubiquitous expression in CNS E18 (RPKM 19.1), adrenal adult (RPKM 18.7) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

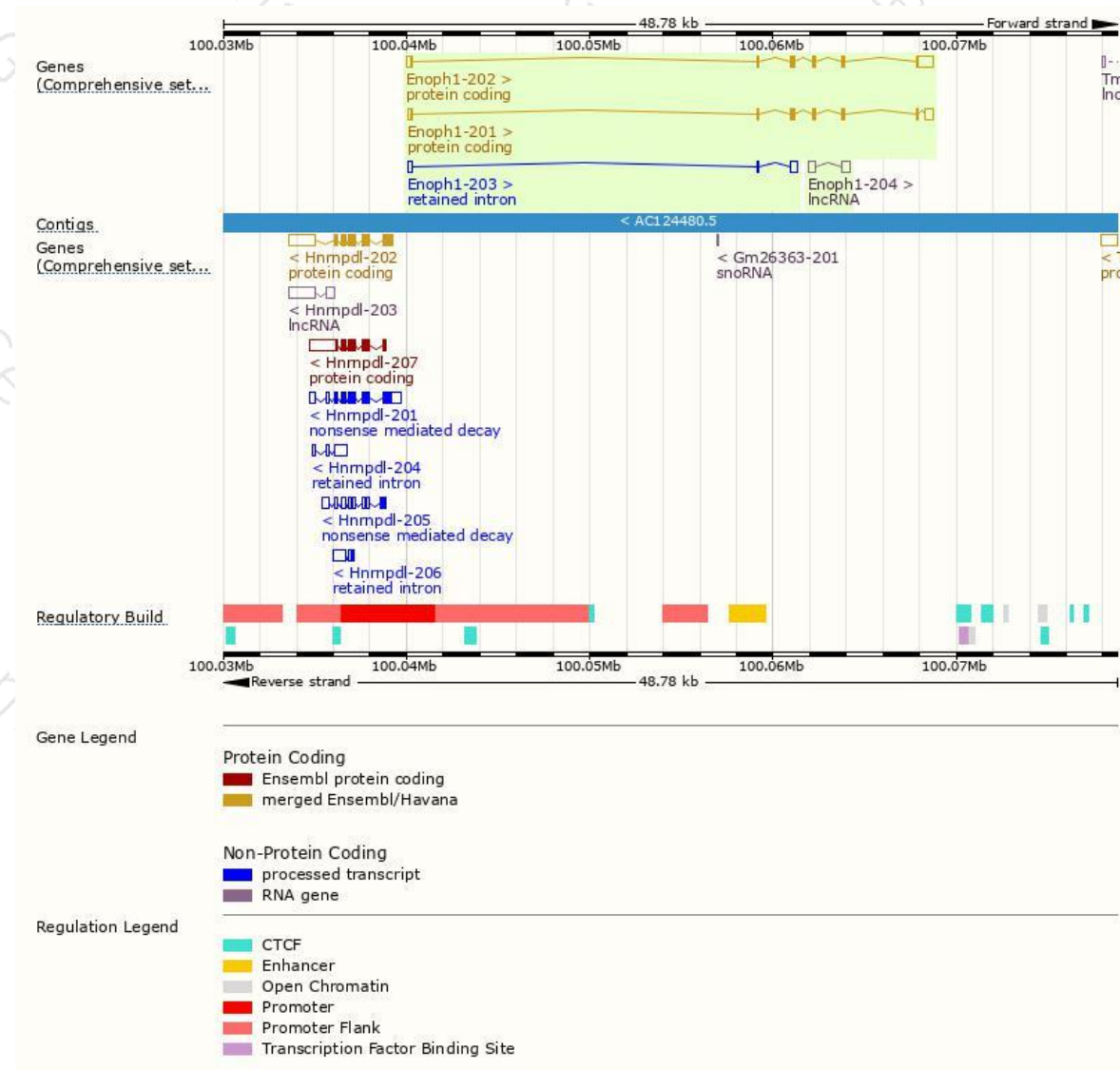
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Enoph1-202	ENSMUST00000169390.7	1833	257aa	Protein coding	CCDS39183	Q8BGB7	TSL:1 GENCODE basic APPRIS P1
Enoph1-201	ENSMUST00000031268.7	1420	257aa	Protein coding	CCDS39183	Q8BGB7	TSL:1 GENCODE basic APPRIS P1
Enoph1-203	ENSMUST00000199518.1	720	No protein	Retained intron	-	-	TSL:1
Enoph1-204	ENSMUST00000200223.1	744	No protein	lncRNA	-	-	TSL:3

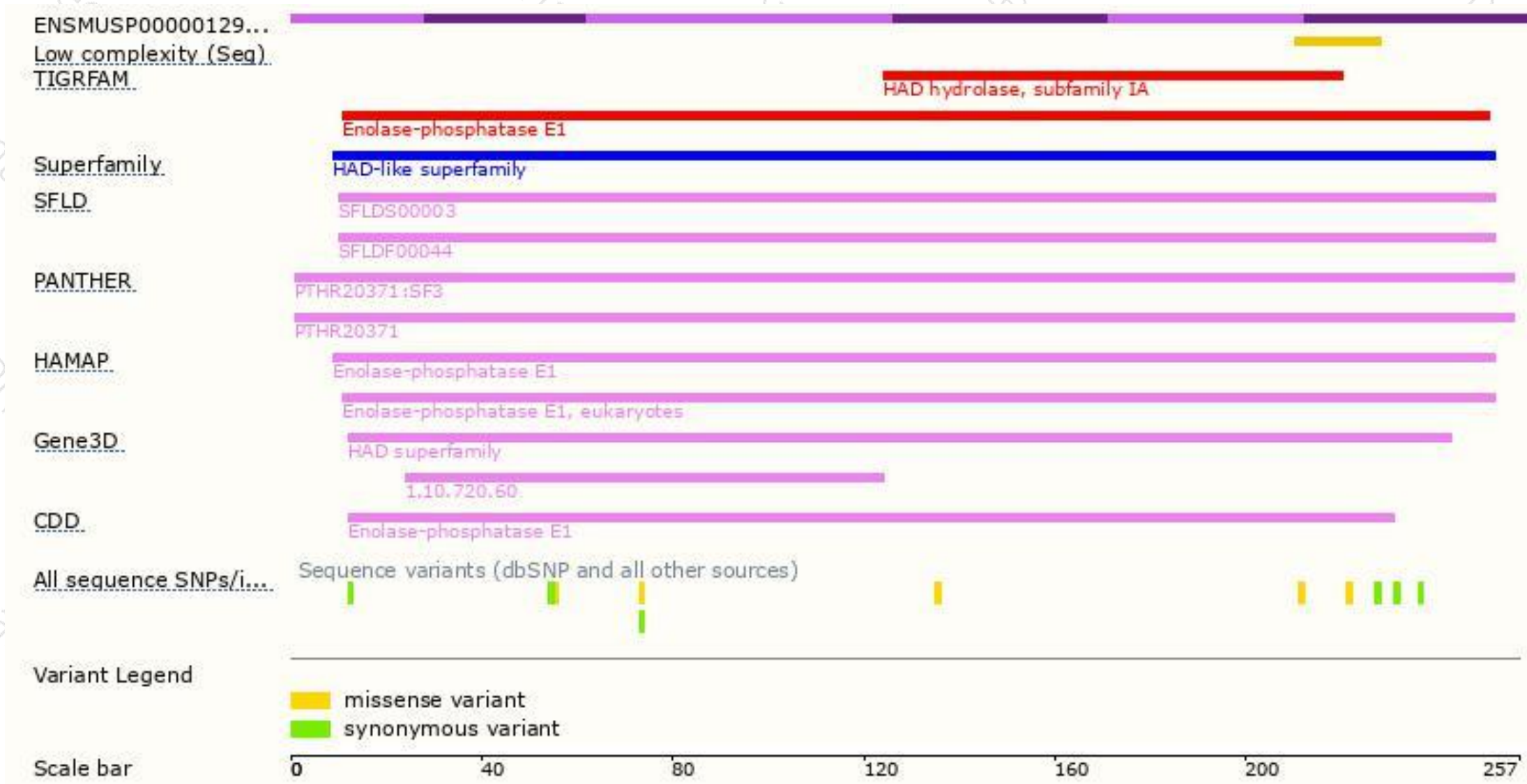
The strategy is based on the design of *Enoph1-202* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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