

***Edn2* Cas9-KO Strategy**

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Project Overview

Project Name

Edn2

Project type

Cas9-KO

Strain background

C57BL/6J

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Edn2* gene. The schematic diagram is as follows:



- The *Edn2* gene has 1 transcript. According to the structure of *Edn2* gene, exon3-exon5 of *Edn2-201* (ENSMUST00000030384.4) transcript is recommended as the knockout region. The region contains 316bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Edn2* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit growth retardation, hypothermia, hypoxemic hypoxia, hypercapnia, emphysema and premature death.
- The *Edn2* gene is located on the Chr4. 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)

Edn2 endothelin 2 [Mus musculus (house mouse)]

Gene ID: 13615, updated on 31-Jan-2019

Summary



Official Symbol Edn2 provided by [MGI](#)

Official Full Name endothelin 2 provided by [MGI](#)

Primary source [MGI:MGI:95284](#)

See related [Ensembl:ENSMUSG00000028635](#)

Gene type protein coding

RefSeq status REVIEWED

Organism [Mus musculus](#)

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as ET-2, PPET2, VIC

Summary This gene encodes a member of the endothelin family of peptides. The encoded preproprotein undergoes proteolytic processing to generate a potent vasoconstrictive peptide. This gene is abundantly expressed in the gastrointestinal tract, strongly induced in photoreceptor cells in retinal diseases and injury, and produced by microglia and macrophages in the early stages of glaucoma. Mice lacking the encoded protein exhibit severe growth retardation, hypothermia and juvenile lethality. [provided by RefSeq, Feb 2016]

Expression Biased expression in large intestine adult (RPKM 14.8), small intestine adult (RPKM 13.6) and 5 other tissues [See more](#)

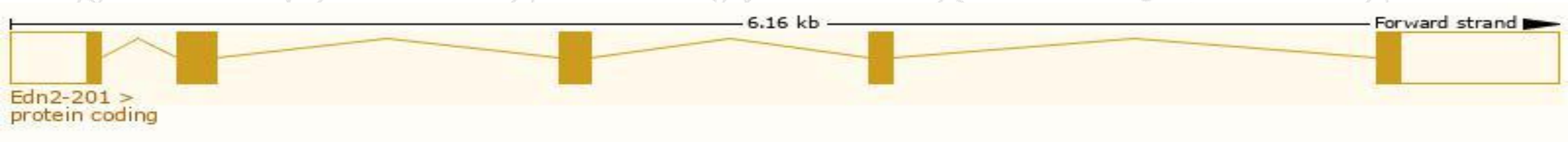
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

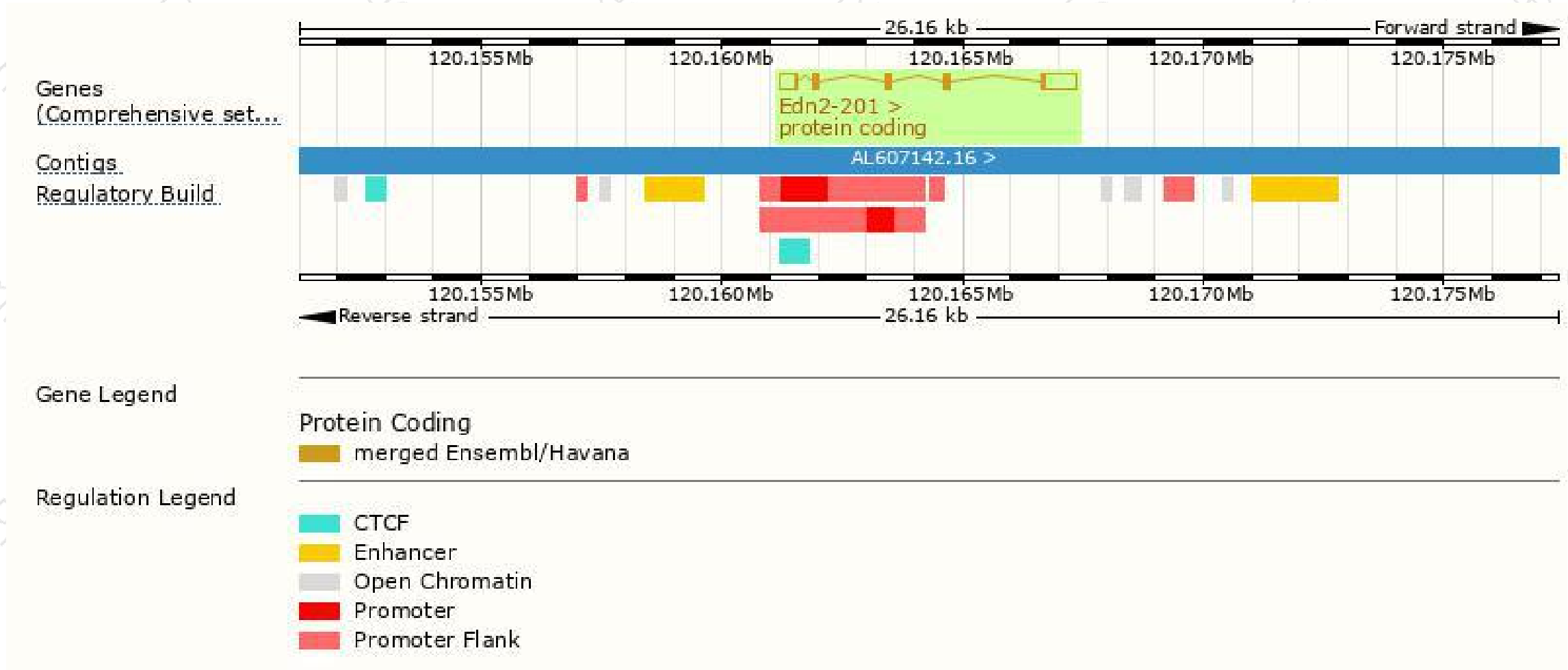
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Edn2-201	ENSMUST00000030384.4	1462	175aa	Protein coding	CCDS38864	P22389	TSL:1 GENCODE basic APPRIS P1

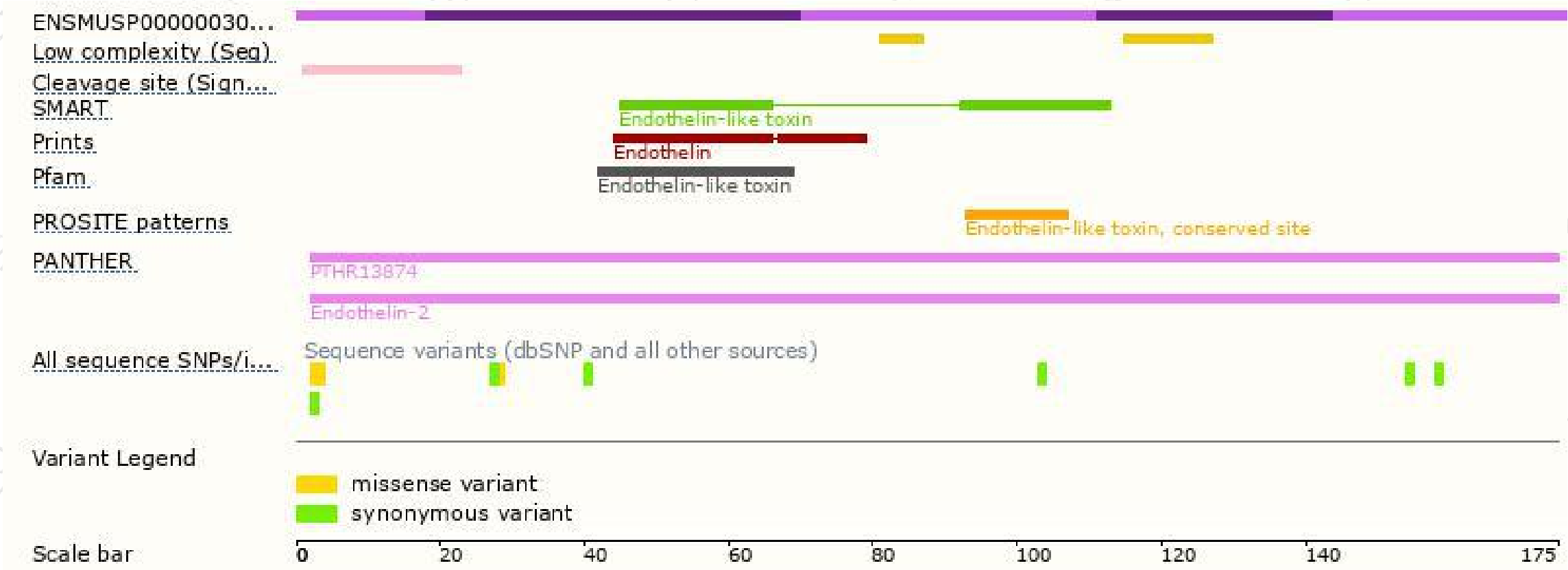
The strategy is based on the design of *Edn2-201* transcript, The transcription is shown below



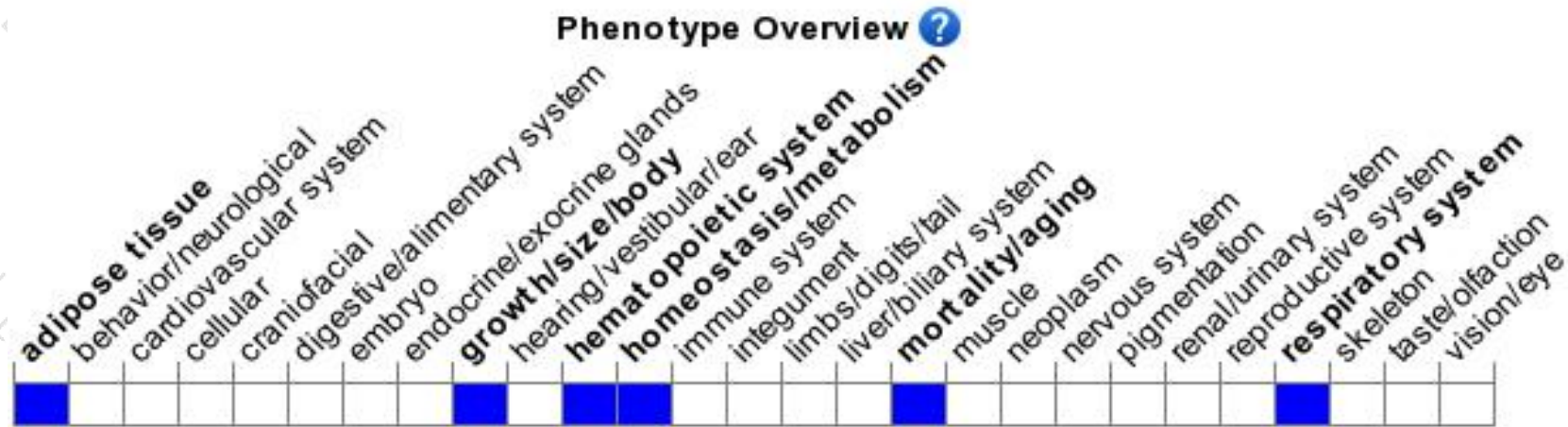
Genomic location distribution



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 exhibit growth retardation, hypothermia, hypoxemic hypoxia, hypercapnia, emphysema and premature death.

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

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