

Notch2 Cas9-KO Strategy

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Project Overview

Project Name

Notch2

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Notch2* gene. The schematic diagram is as follows:



- The *Notch2* gene has 2 transcripts. According to the structure of *Notch2* gene, exon2 of *Notch2-201* (ENSMUST00000079812.7) transcript is recommended as the knockout region. The region contains 82bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Notch2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygotes for null alleles exhibit defects in embryonic development resulting in embryonic or neonatal lethality.
- The *Notch2* gene is located on the Chr3. 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)

Notch2 notch 2 [Mus musculus (house mouse)]

Gene ID: 18129, updated on 2-Apr-2019

Summary



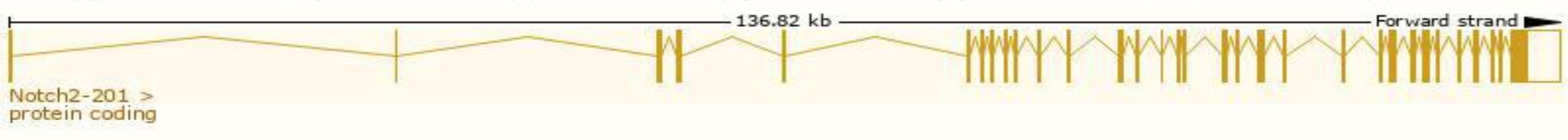
Official Symbol	Notch2 provided by MGI
Official Full Name	notch 2 provided by MGI
Primary source	MGI:MGI:97364
See related	Ensembl:ENSMUSG00000027878
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI853703, N2
Expression	Ubiquitous expression in spleen adult (RPKM 26.6), ovary adult (RPKM 22.8) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

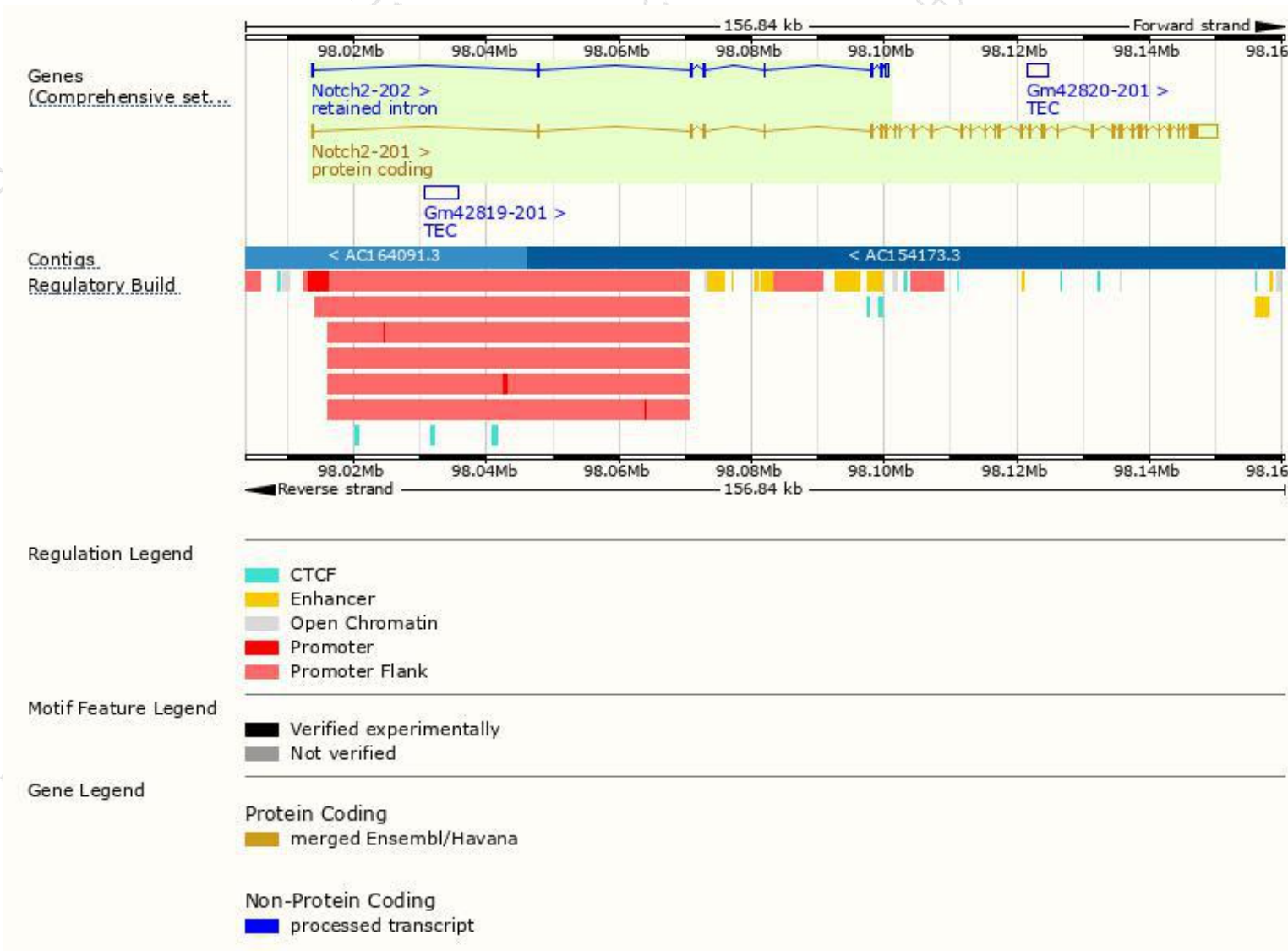
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Notch2-201	ENSMUST00000079812.7	10500	2473aa	Protein coding	CCDS51013	G5E8J0	TSL:1 GENCODE basic APPRIS P1
Notch2-202	ENSMUST00000198324.1	1896	No protein	Retained intron	-	-	TSL:1

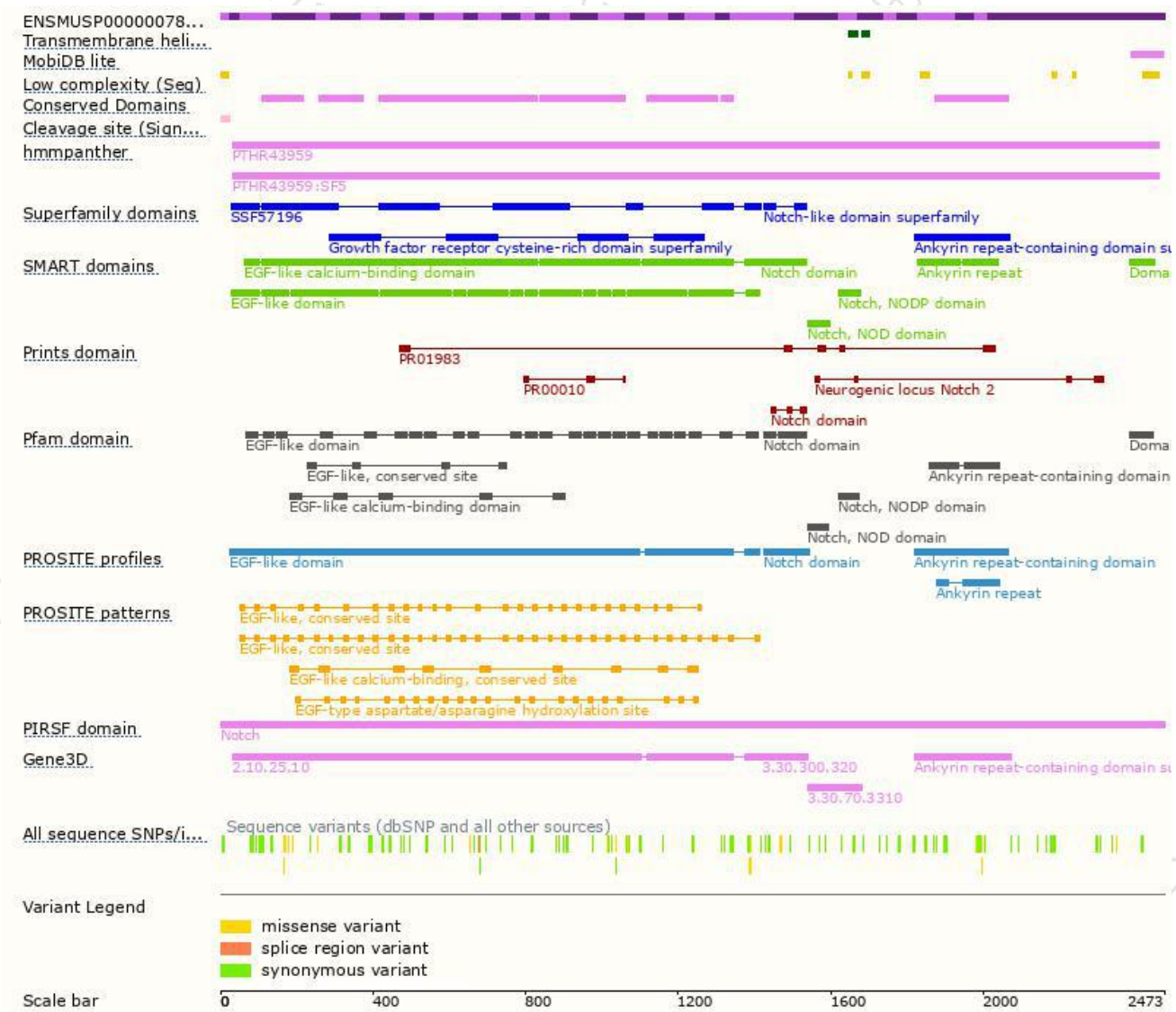
The strategy is based on the design of *Notch2-201* transcript,The transcription is shown below



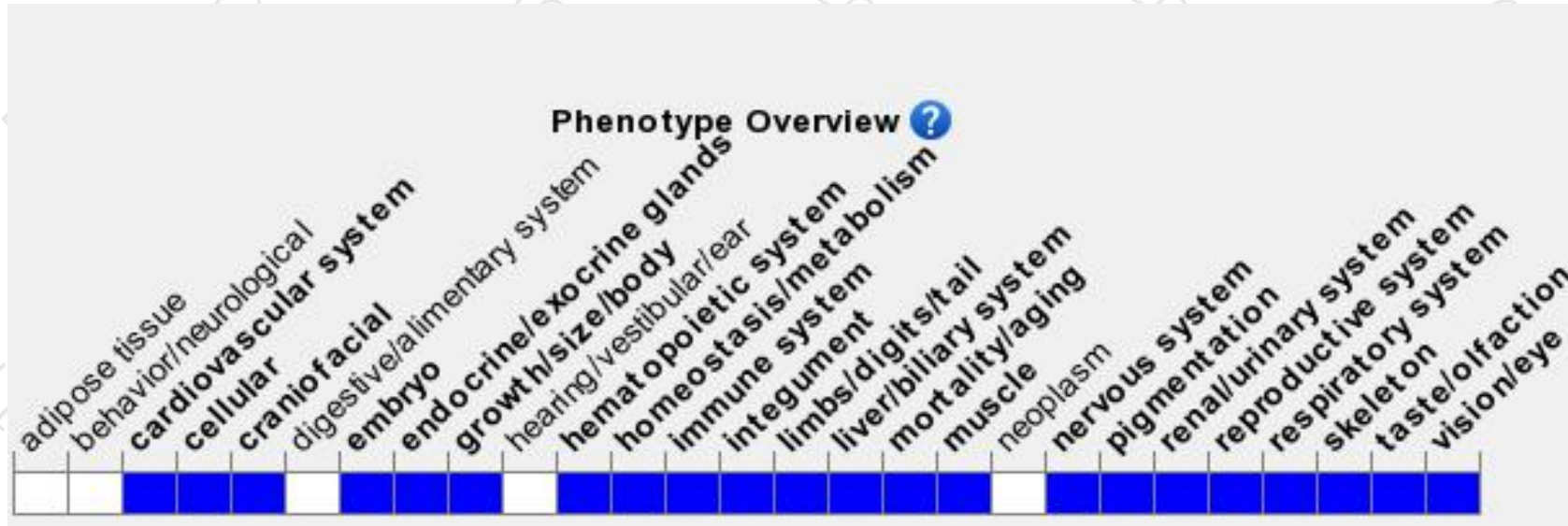
Genomic location distribution



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, Homozygotes for null alleles exhibit defects in embryonic development resulting in embryonic or neonatal lethality.

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

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