

Rdh1 Cas9-KO Strategy

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Reviewer:

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Design Date:

2020-2-24

Project Overview

Project Name

Rdh1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rdh1* gene. The schematic diagram is as follows:



- The *Rdh1* gene has 1 transcript. According to the structure of *Rdh1* gene, exon2 of *Rdh1-201* (ENSMUST00000073639.5) transcript is recommended as the knockout region. The region contains 259bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rdh1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Mice homozygous for a null allele exhibit increased body weight, adipose tissue, and retinol levels.
- The flox region is in the intron of the *Rdh16f1* gene, which may affect the regulation of this gene.
- The *Rdh1* gene is located on the Chr10. 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)

Rdh1 retinol dehydrogenase 1 (all trans) [Mus musculus (house mouse)]

Gene ID: 107605, updated on 19-Mar-2019

Summary

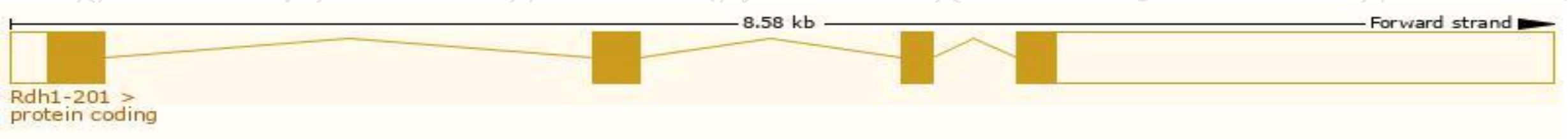
Official Symbol	Rdh1 provided by MGI
Official Full Name	retinol dehydrogenase 1 (all trans) provided by MGI
Primary source	MGI:MGI:1195275
See related	Ensembl:ENSMUSG000000089789
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Biased expression in liver adult (RPKM 10.7), liver E18 (RPKM 2.1) and 4 other tissues See more

Transcript information (Ensembl)

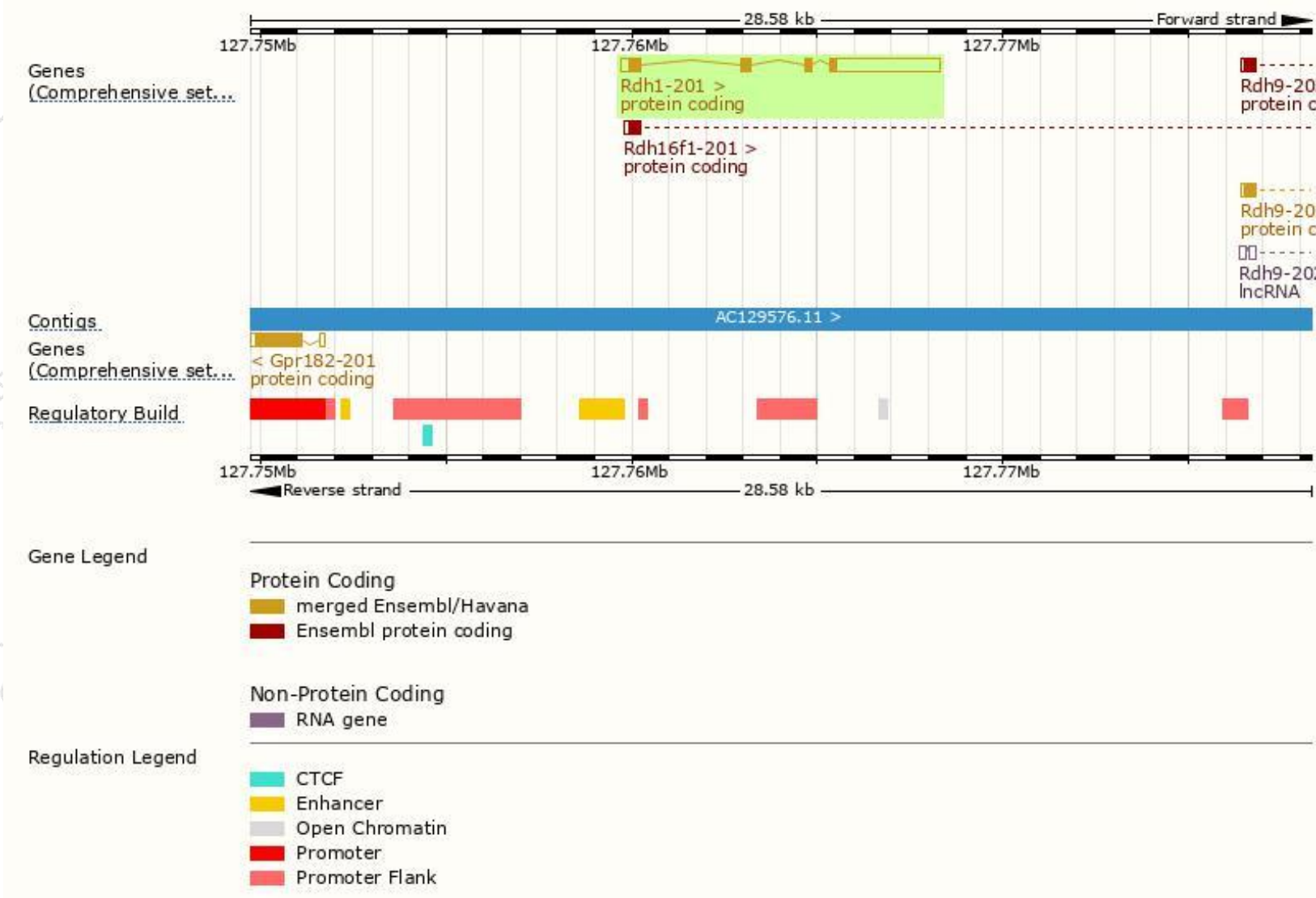
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rdh1-201	ENSMUST00000073639.5	3930	317aa	Protein coding	CCDS24250	Q8CGV4	TSL:1 GENCODE basic APPRIS P1

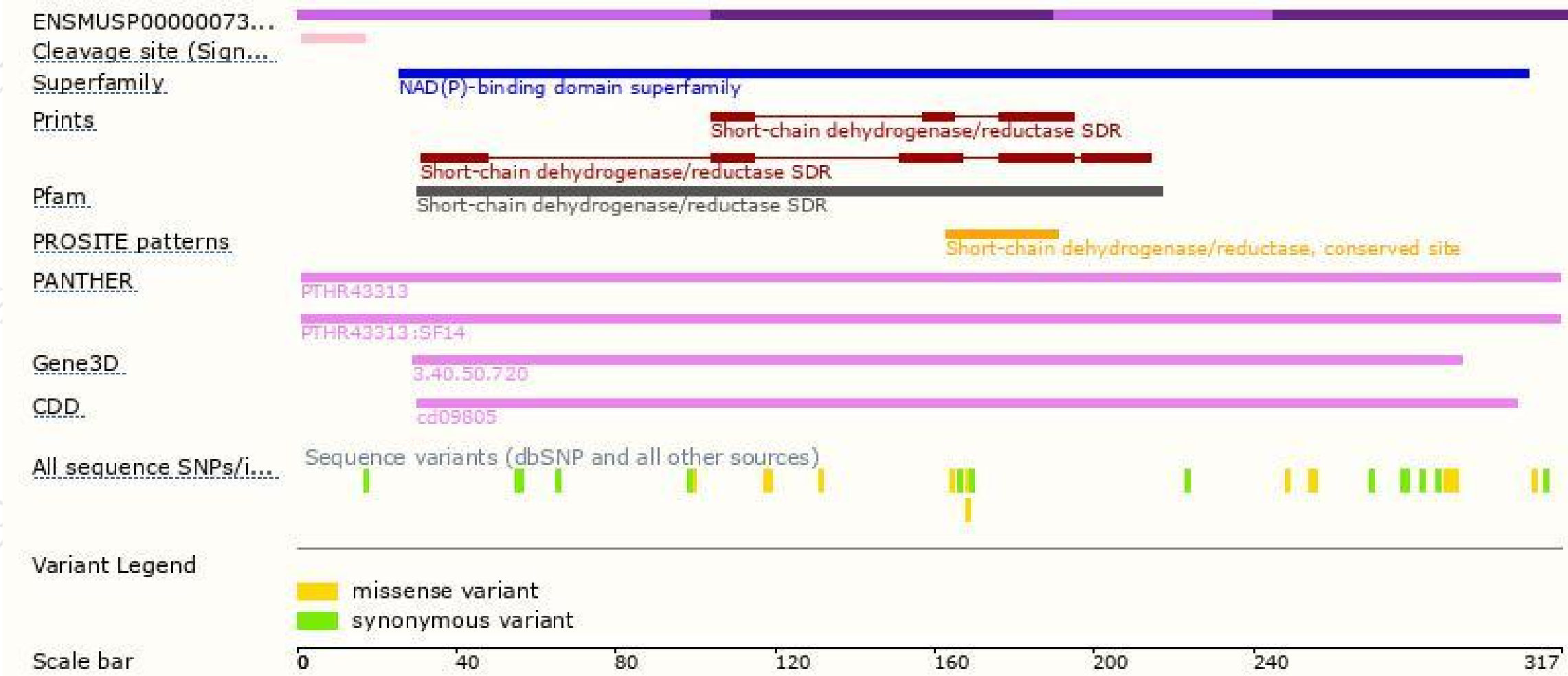
The strategy is based on the design of *Rdh1-201* transcript, The transcription is shown below



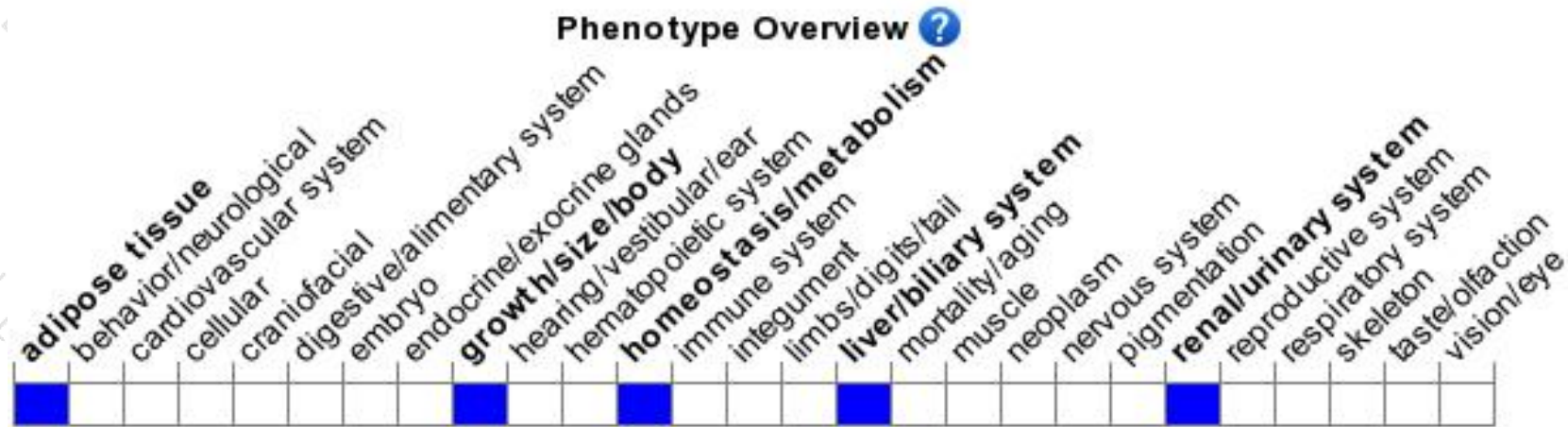
Genomic location distribution



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 null allele exhibit increased body weight, adipose tissue, and retinol levels.

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

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