

***Rbp1* Cas9-KO Strategy**

Designer: Xueting Zhang

Reviewer: Yanhua Shen

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Project Overview

Project Name

Rbp1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rbp1* gene. The schematic diagram is as follows:



- The *Rbp1* gene has 1 transcript. According to the structure of *Rbp1* gene, exon1-exon4 of *Rbp1-201* (ENSMUST00000052068.10) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rbp1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Homozygotes for a targeted null mutation show increased susceptibility to a diet deficient in vitamin A. Mutants also exhibit a two-fold delay in dark adaptation after a photic flash.
- The *Rbp1* gene is located on the Chr9. 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)

Rbp1 retinol binding protein 1, cellular [*Mus musculus* (house mouse)]

Gene ID: 19659, updated on 12-Aug-2019

Summary

Official Symbol Rbp1 provided by [MGI](#)
Official Full Name retinol binding protein 1, cellular provided by [MGI](#)
Primary source [MGI:MGI:97876](#)
See related [Ensembl:ENSMUSG00000046402](#)
Gene type protein coding
RefSeq status PROVISIONAL
Organism [Mus musculus](#)
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as Crbp; CRBPI; Rbp-1
Expression Broad expression in placenta adult (RPKM 52.2), liver E18 (RPKM 35.5) and 18 other tissues [See more](#)
Orthologs [human](#) [all](#)

Genomic context

Location: 9 E3.3; 9 51.36 cM

See Rbp1 in [Genome Data Viewer](#)

Exon count: 4

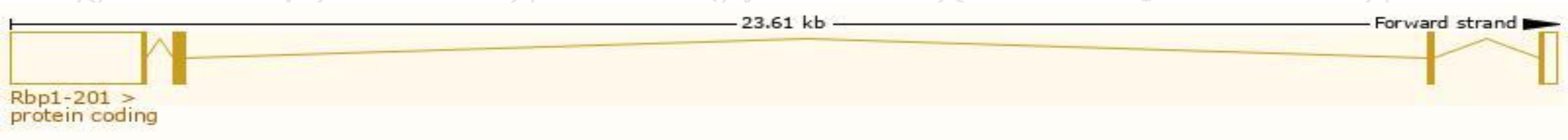
| Annotation release | Status | Assembly | Chr | Location |
|---------------------|-------------------|--|-----|----------------------------------|
| 108 | current | GRCm38.p6 (GCF_000001635.26) | 9 | NC_000075.6 (98422961..98446551) |
| Build 37.2 | previous assembly | MGSCv37 (GCF_000001635.18) | 9 | NC_000075.5 (98323380..98346970) |

Transcript information (Ensembl)

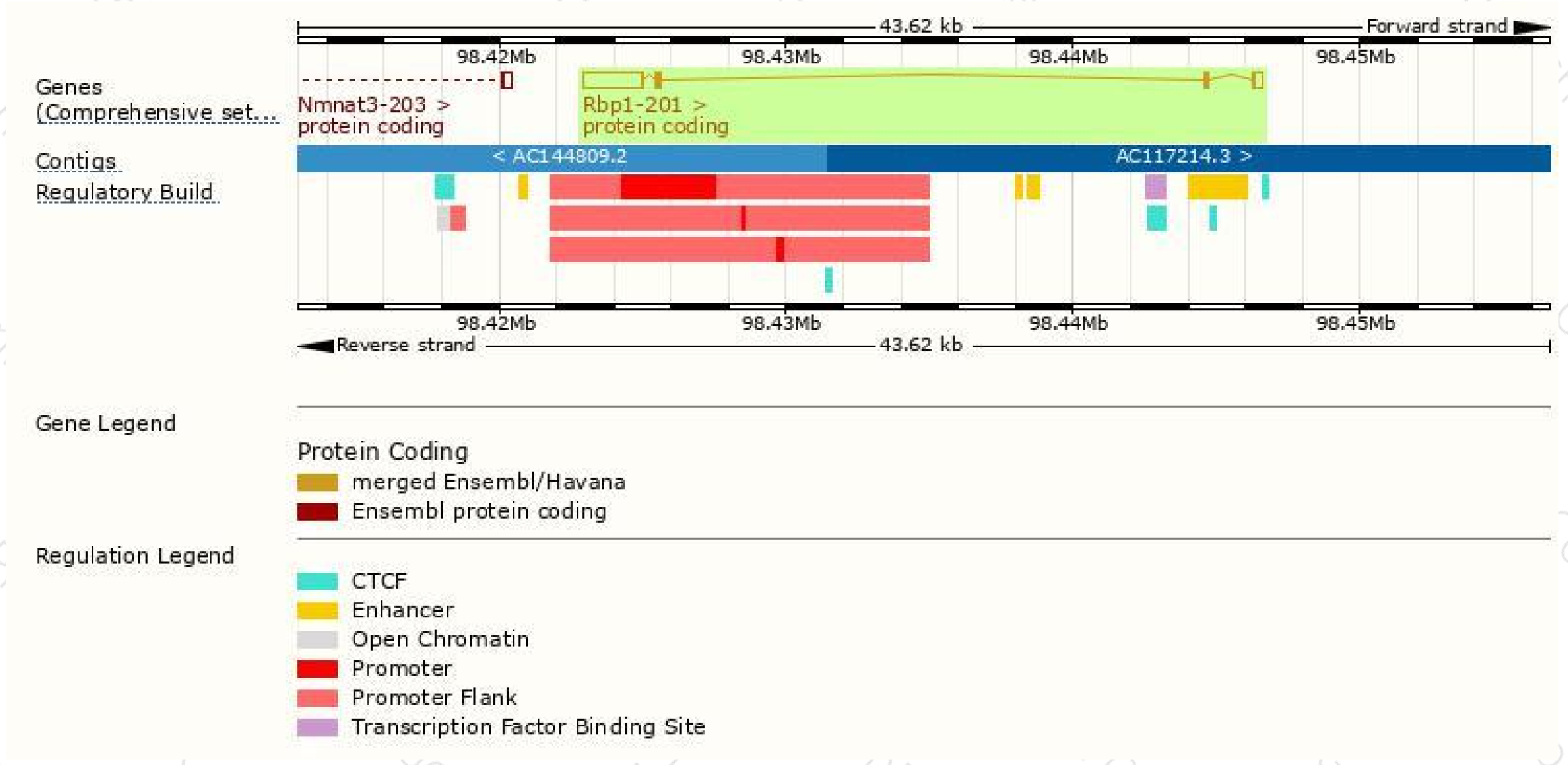
The gene has 1 transcript, and the transcript is shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|----------|---------------------------------------|------|-----------------------|----------------|---------------------------|-------------------------------|-------------------------------|
| Rbp1-201 | ENSMUST00000052068.10 | 2652 | 135aa | Protein coding | CCDS23424 | Q00915 Q58EU7 | TSL:1 GENCODE basic APPRIS P1 |

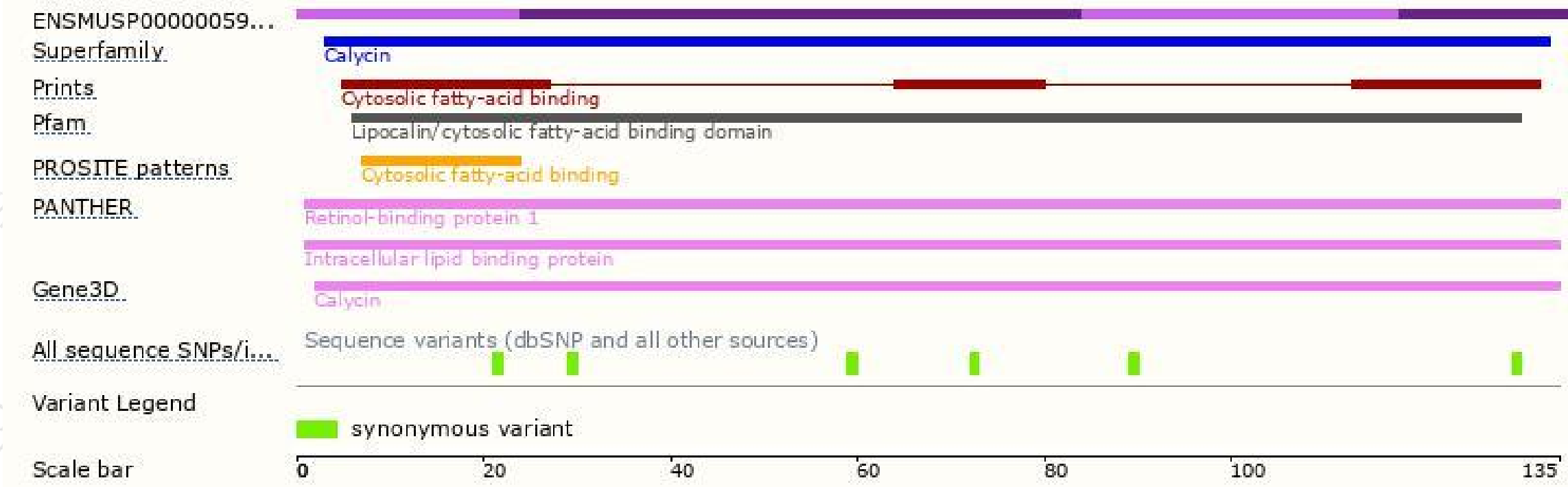
The strategy is based on the design of *Rbp1-201* transcript, The transcription is shown below



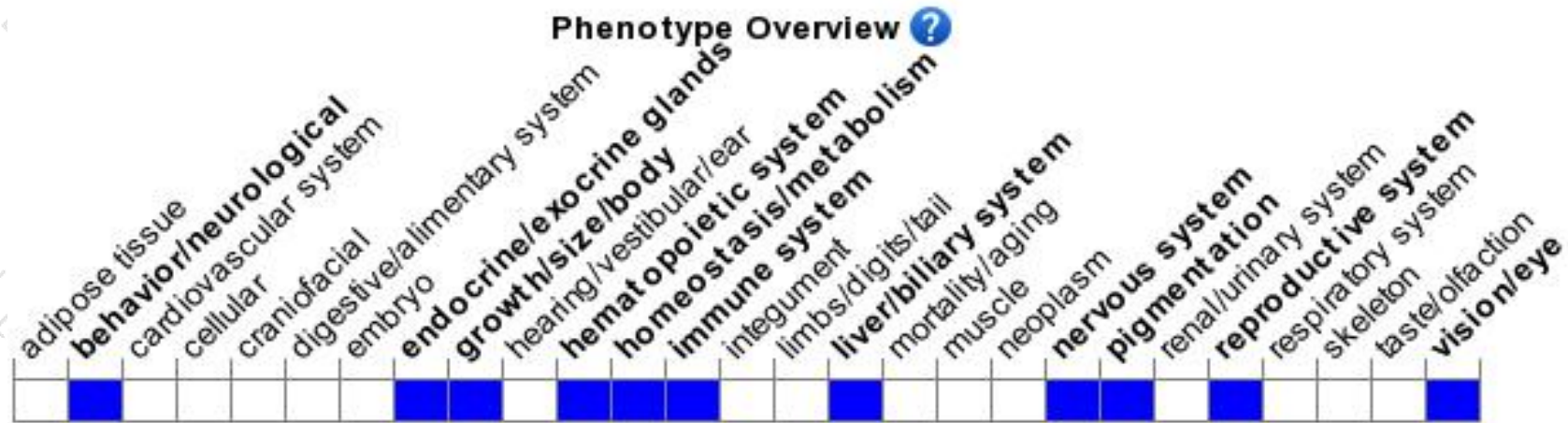
Genomic location distribution



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 a targeted null mutation show increased susceptibility to a diet deficient in vitamin A. Mutants also exhibit a two-fold delay in dark adaptation after a photic flash.

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

Tel: 400-9660890

