

Orm1 Cas9-KO Strategy

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Project Overview

Project Name

Orm1

Project type

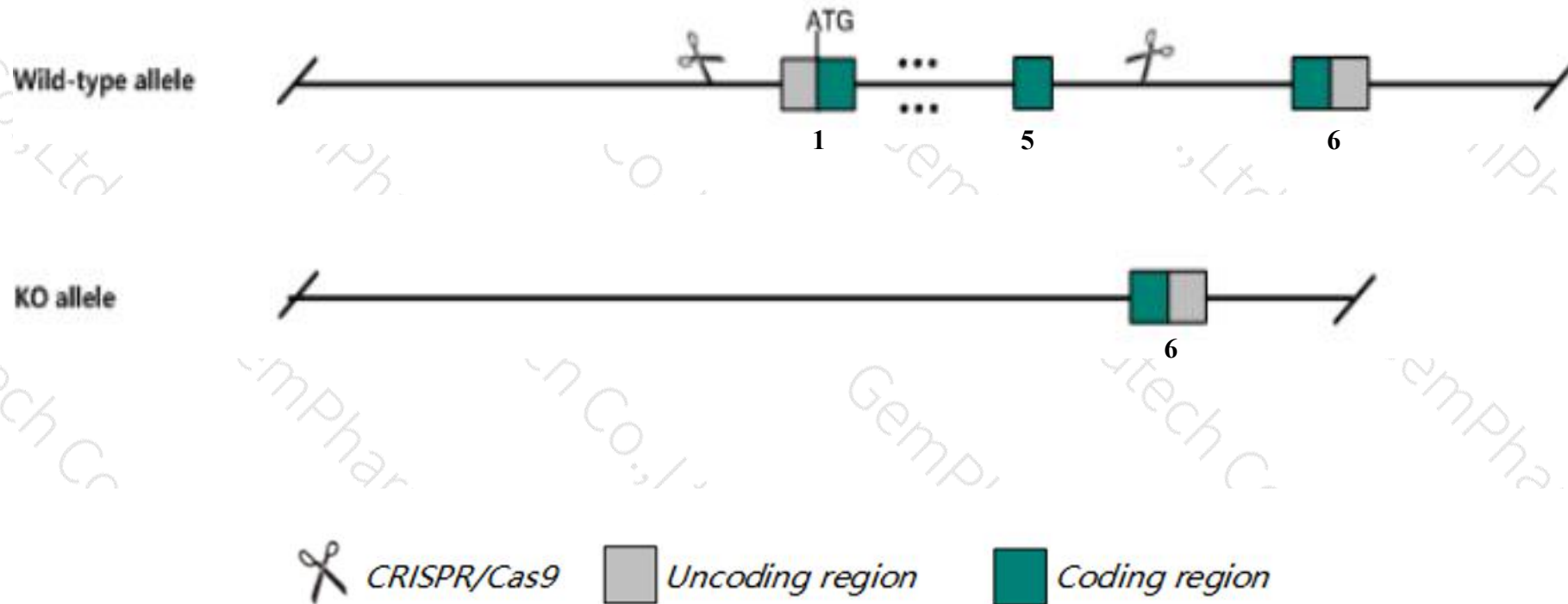
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Orm1* gene. The schematic diagram is as follows:



- The *Orm1* gene has 1 transcript. According to the structure of *Orm1* gene, exon1-exon5 of *Orm1-201* (ENSMUST00000030044.2) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Orm1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Orm1* 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)

Orm1 orosomucoid 1 [Mus musculus (house mouse)]

Gene ID: 18405, updated on 13-Mar-2020

Summary



Official Symbol Orm1 provided by [MGI](#)

Official Full Name orosomucoid 1 provided by [MGI](#)

Primary source [MGI:MGI:97443](#)

See related [Ensembl:ENSMUSG00000039196](#)

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 Agp-1, Agp-2, Orm-1

Summary This gene encodes a member of the lipocalin family of proteins. The encoded protein is an abundant acute-phase protein that is synthesized by hepatocytes in response to cytokines during the acute phase response. The encoded protein may regulate inflammation and metabolism. This gene is present in a gene cluster on chromosome 4. [provided by RefSeq, Aug 2015]

Expression Biased expression in liver adult (RPKM 234.2), subcutaneous fat pad adult (RPKM 106.0) and 2 other tissues [See more](#)

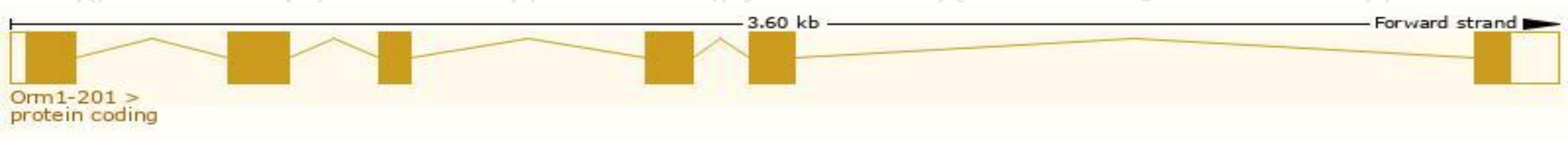
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

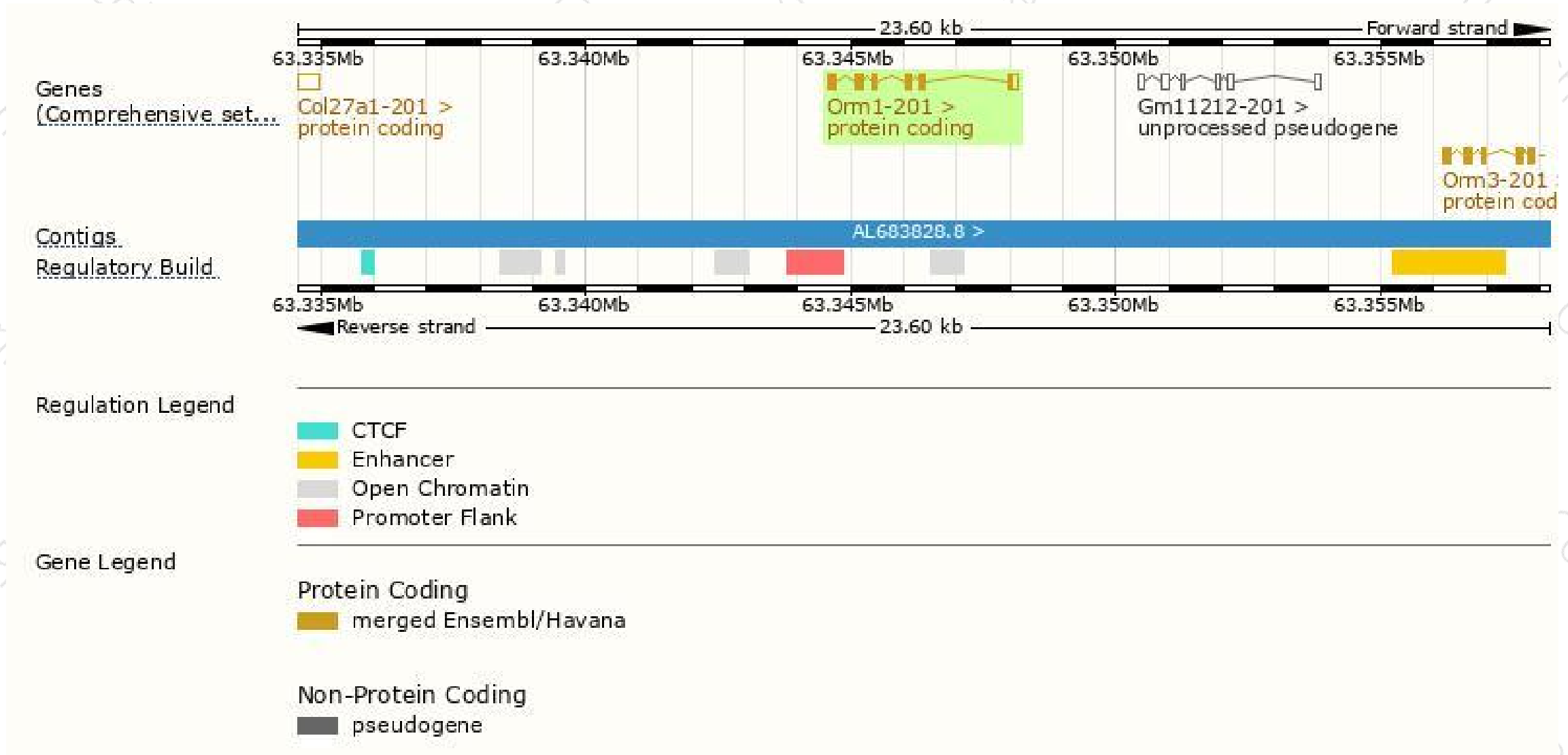
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Orm1-201	ENSMUST00000030044.2	775	207aa	Protein coding	CCDS18251	Q60590	TSL:1 GENCODE basic APPRIS P1

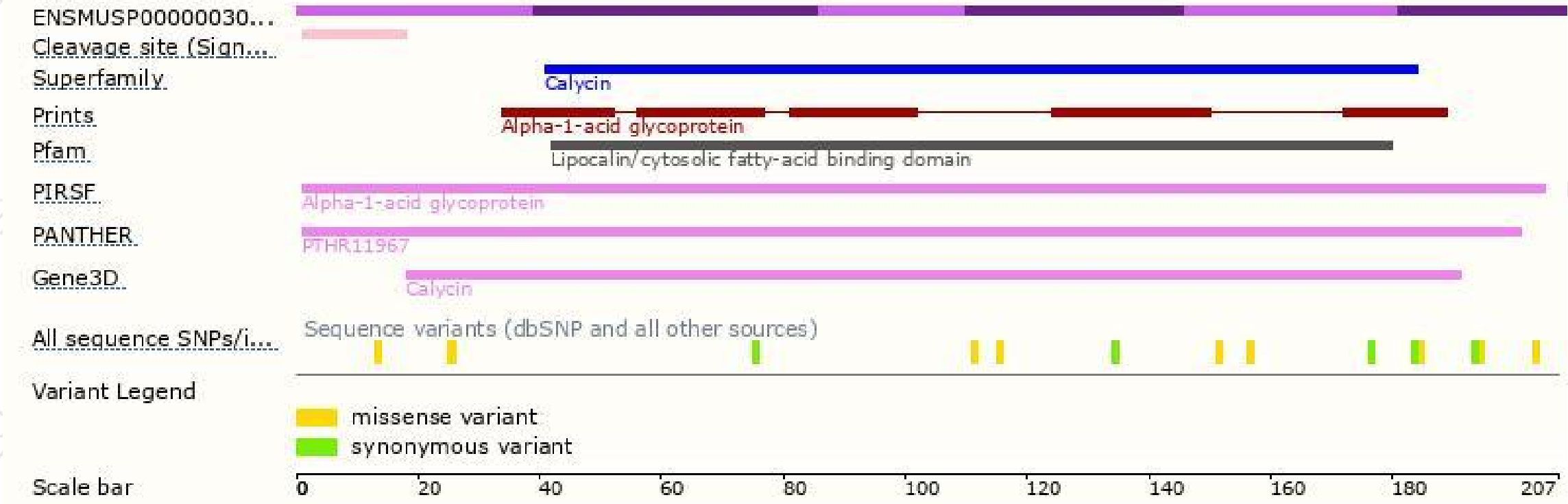
The strategy is based on the design of *Orm1-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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