

C1ql2 Cas9-KO Strategy

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Project Overview

Project Name

C1ql2

Project type

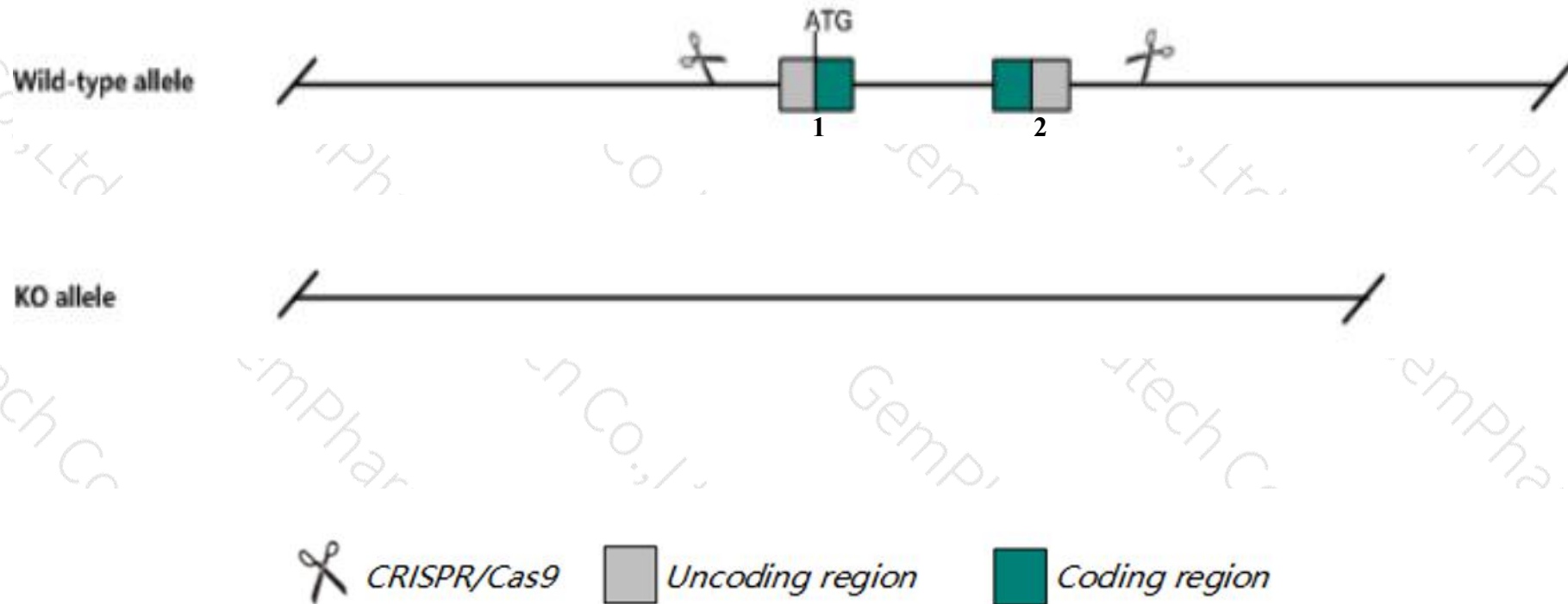
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Clql2* gene. The schematic diagram is as follows:



- The *Clql2* gene has 1 transcript. According to the structure of *Clql2* gene, exon1-exon2 of *Clql2-201* (ENSMUST00000037286.9) 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 *Clql2* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Clql2* gene is located on the Chr1. 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)

C1ql2 complement component 1, q subcomponent-like 2 [Mus musculus (house mouse)]

Gene ID: 226359, updated on 13-Mar-2020

Summary



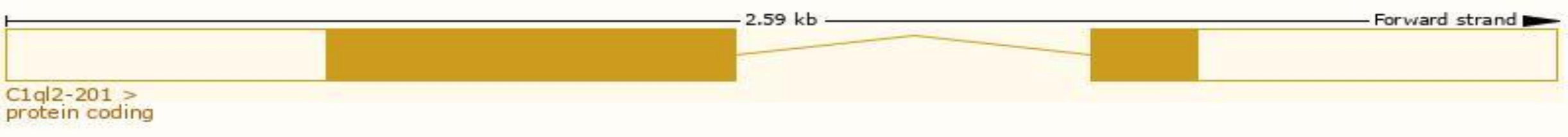
Official Symbol	C1ql2 provided by MGI
Official Full Name	complement component 1, q subcomponent-like 2 provided by MGI
Primary source	MGI:MGI:3032521
See related	Ensembl:ENSMUSG00000036907
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Aii, BC040774, CTRP10
Expression	Biased expression in cortex adult (RPKM 1.9), CNS E18 (RPKM 1.8) and 8 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

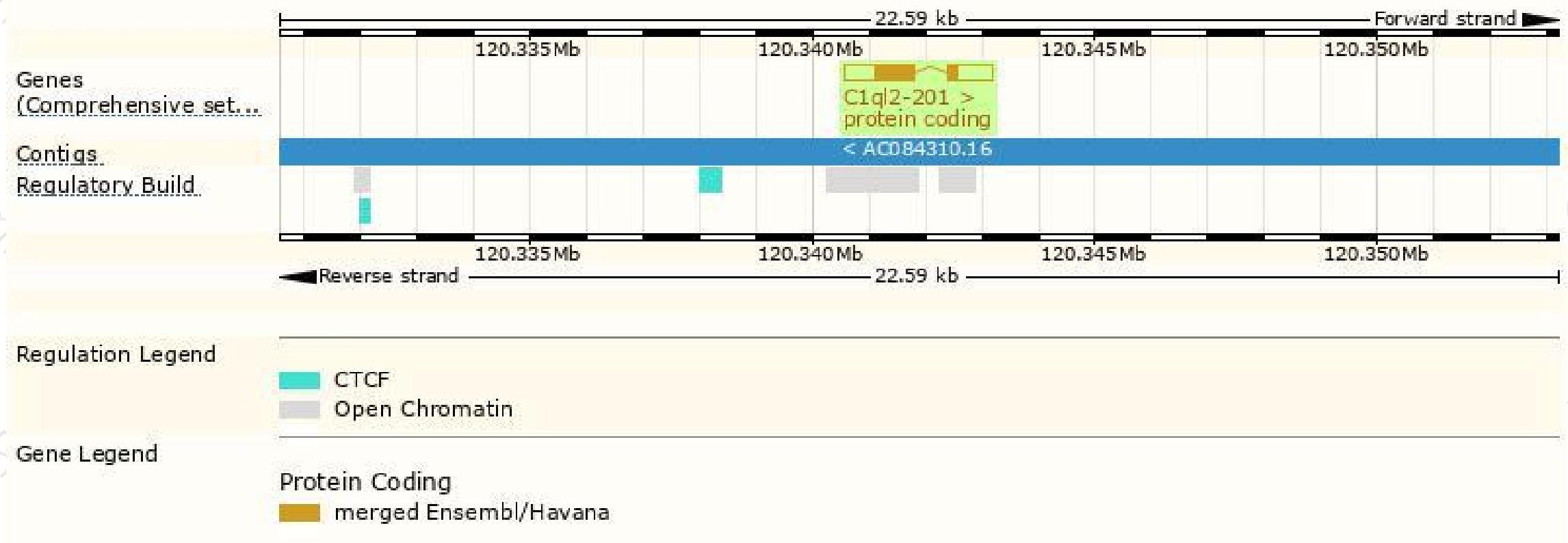
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
C1ql2-201	ENSMUST00000037286.9	2000	287aa	Protein coding	CCDS15233	A0A3B0J351_Q8CFR0	TSL:1 GENCODE basic APPRIS P1

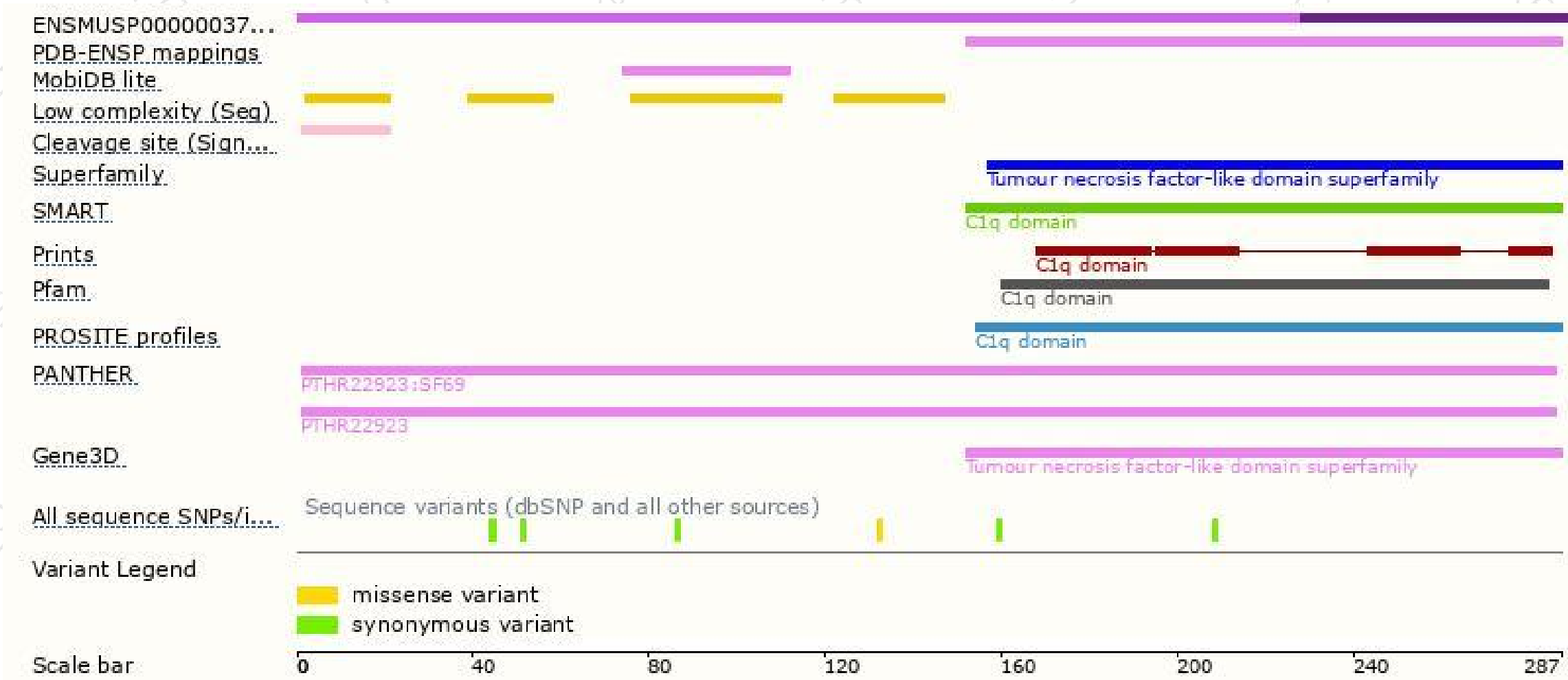
The strategy is based on the design of *C1ql2-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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