

Yaf2 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Yaf2

Project type

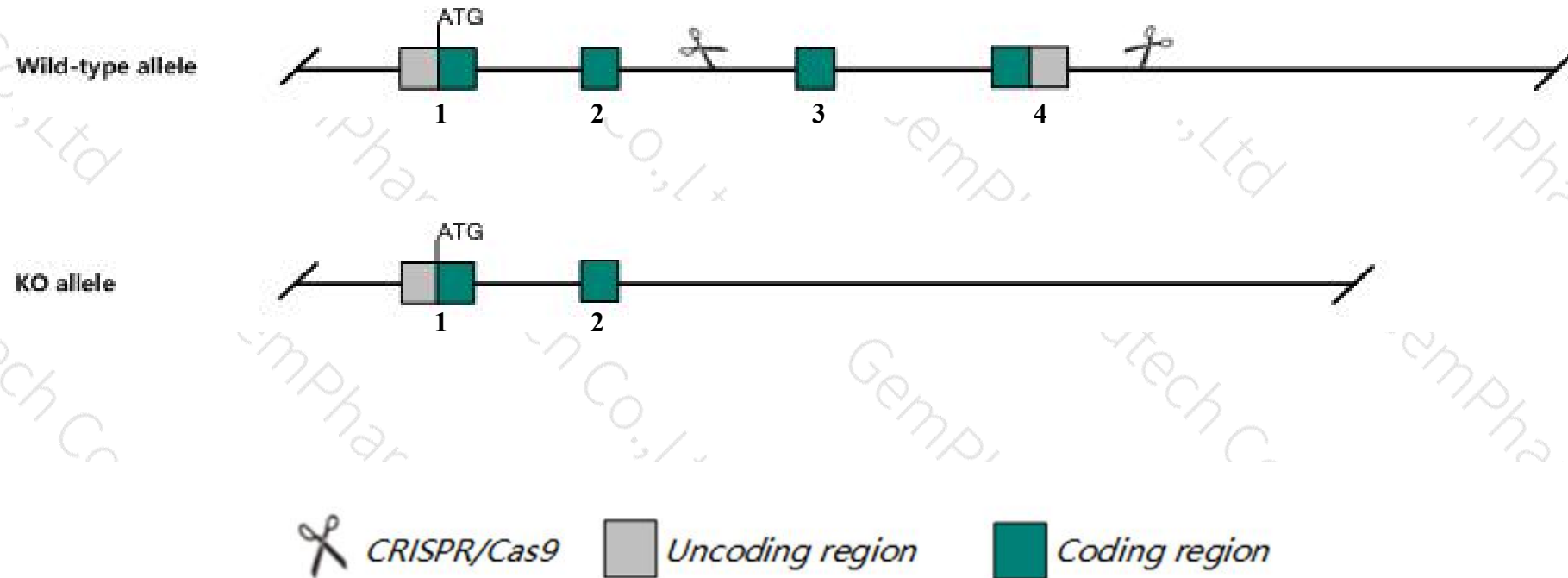
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Yaf2* gene. The schematic diagram is as follows:



- The *Yaf2* gene has 2 transcripts. According to the structure of *Yaf2* gene, exon3-exon4 of *Yaf2-201* (ENSMUST00000080299.6) transcript is recommended as the knockout region. The region contains 388bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Yaf2* gene. The brief process is as follows: CRISPR/Cas9 system w

- The *Yaf2* gene is located on the Chr15. 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)

Yaf2 YY1 associated factor 2 [Mus musculus (house mouse)]

Gene ID: 67057, updated on 31-Jan-2019

Summary



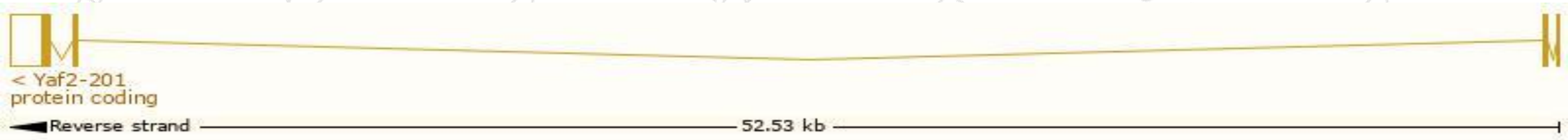
Official Symbol	Yaf2 provided by MGI
Official Full Name	YY1 associated factor 2 provided by MGI
Primary source	MGI:MGI:1914307
See related	Ensembl:ENSMUSG00000022634
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2810021M11Rik
Expression	Ubiquitous expression in cortex adult (RPKM 27.2), frontal lobe adult (RPKM 27.0) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

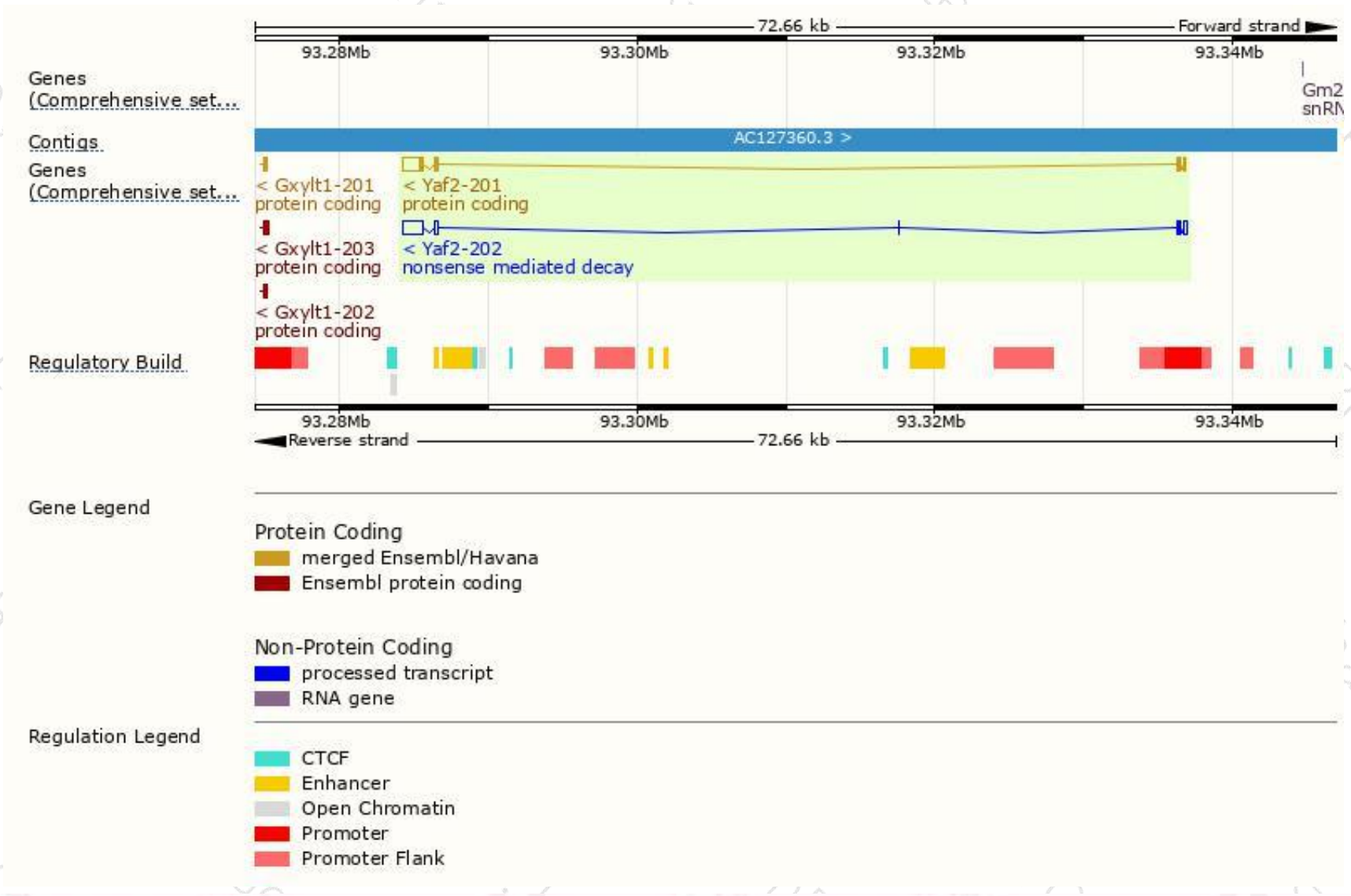
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Yaf2-201	ENSMUST00000080299.6	1676	179aa	Protein coding	CCDS27765	Q99LW6	TSL:1 GENCODE basic APPRIS P1
Yaf2-202	ENSMUST00000133736.1	1870	68aa	Nonsense mediated decay	-	Q99LW6	TSL:1

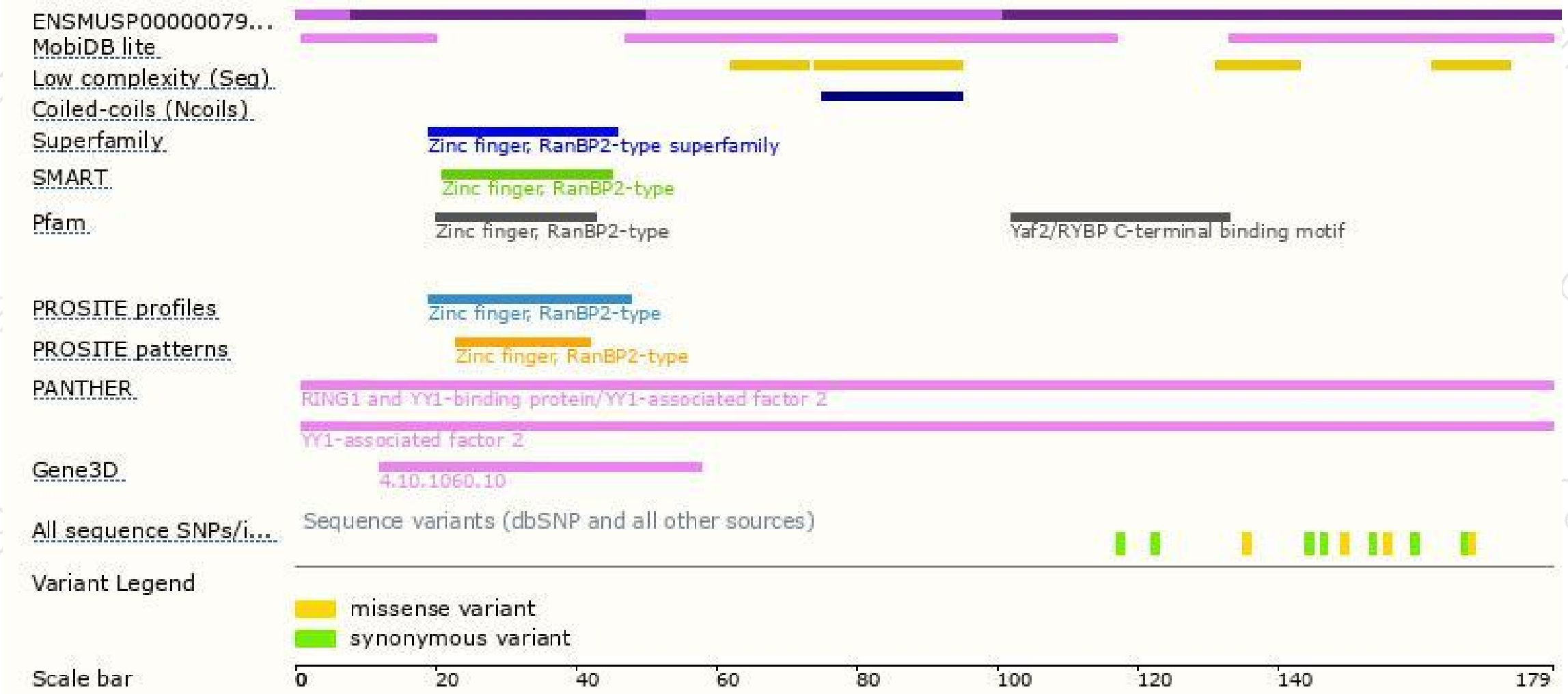
The strategy is based on the design of *Yaf2-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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