

***Rell1* Cas9-KO Strategy**

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Project Overview

Project Name

Rel1

Project type

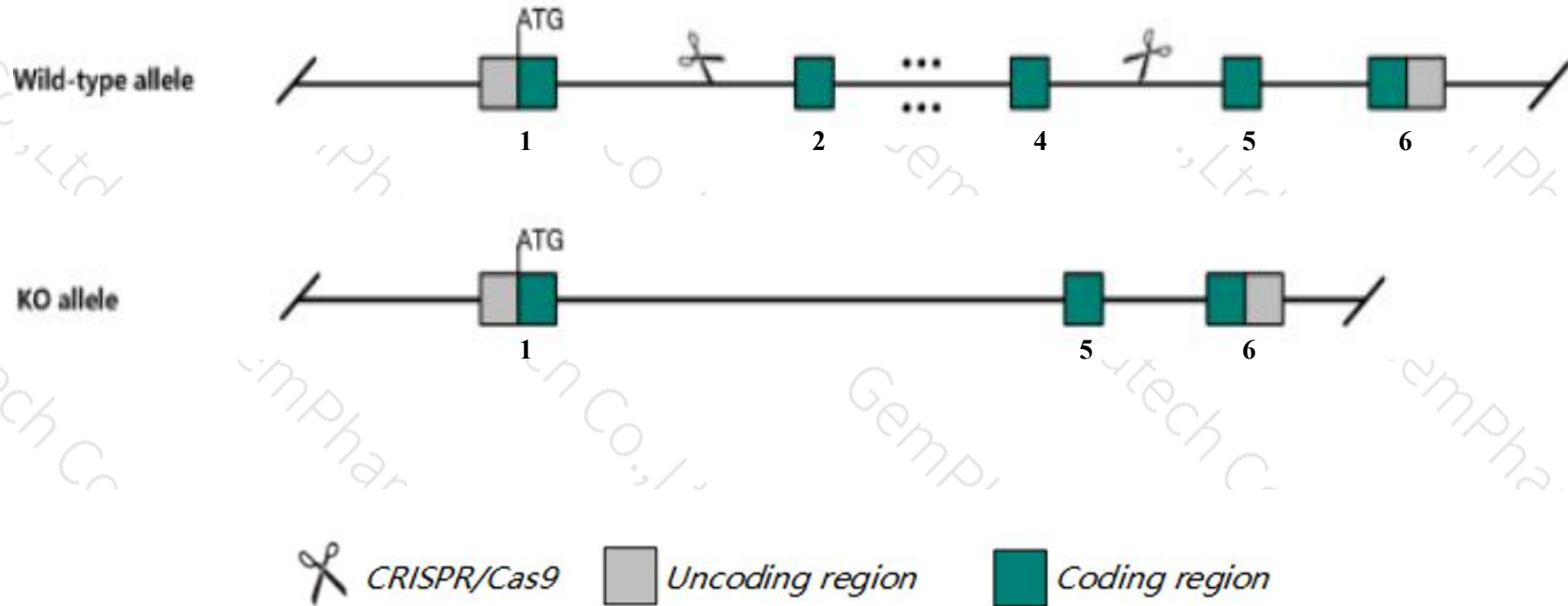
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rel1* gene. The schematic diagram is as follows:



- The *Rel1* gene has 5 transcripts. According to the structure of *Rel1* gene, exon2-exon4 of *Rel1-204* (ENSMUST00000154169.3) transcript is recommended as the knockout region. The region contains 358bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rel1* gene. The brief process is as follows: CRISPR/Cas9 system v

- The *Rel1* gene is located on the Chr5. 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)

Rel1 RELT-like 1 [Mus musculus (house mouse)]

Gene ID: 100532, updated on 13-Mar-2020

Summary



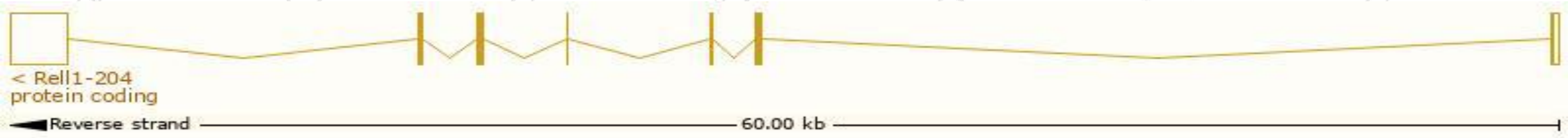
Official Symbol	Rel1 provided by MGI
Official Full Name	RELT-like 1 provided by MGI
Primary source	MGI:MGI:2140767
See related	Ensembl:ENSMUSG00000047881
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AA536743
Expression	Ubiquitous expression in placenta adult (RPKM 8.9), adrenal adult (RPKM 7.4) and 27 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

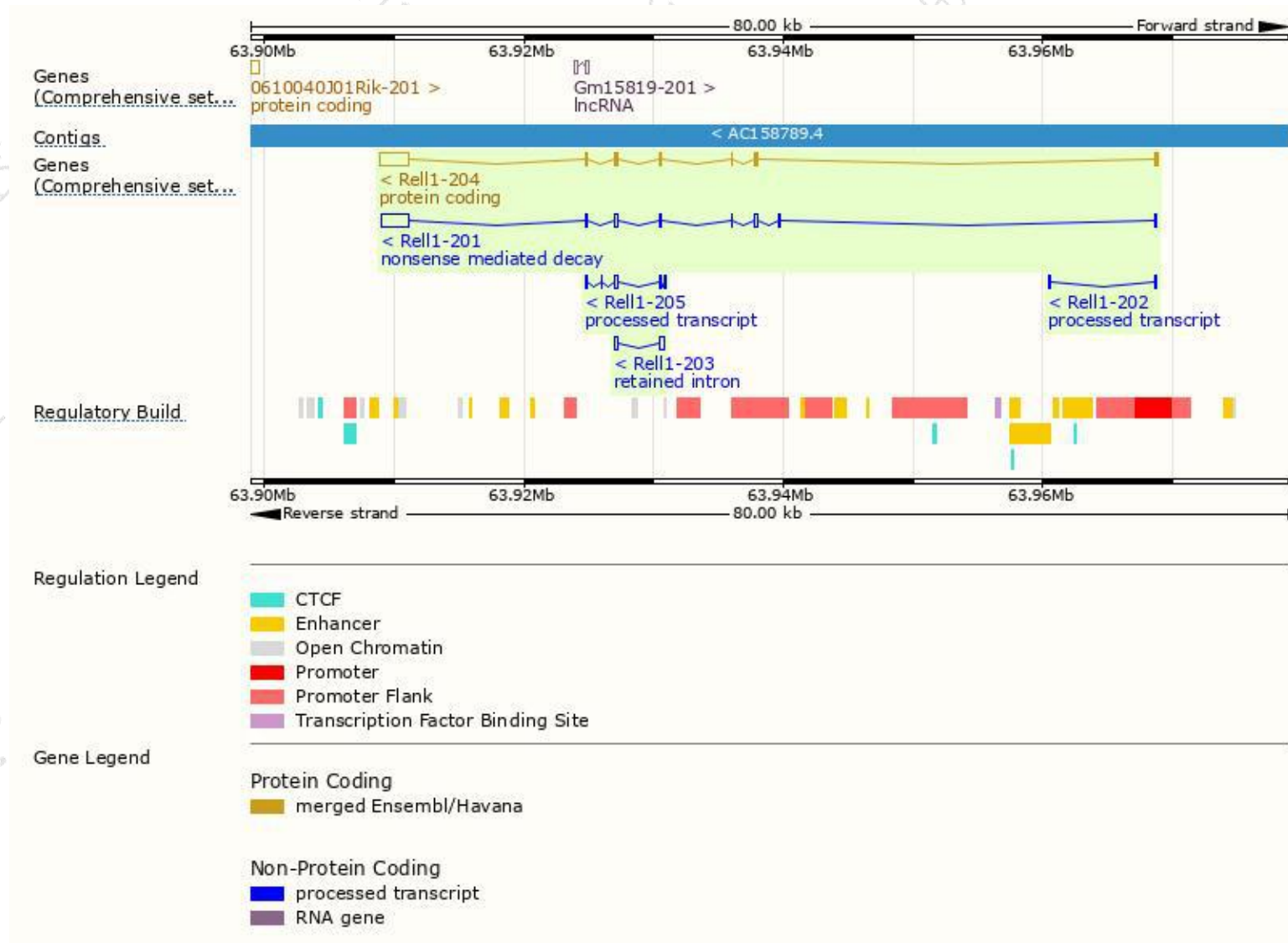
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rel1-204	ENSMUST00000154169.3	3241	272aa	Protein coding	CCDS39093	Q8K2J7	TSL:1 GENCODE basic APPRIS P1
Rel1-201	ENSMUST00000087327.12	3093	57aa	Nonsense mediated decay	-	H7BX61	TSL:1
Rel1-205	ENSMUST00000154532.2	490	No protein	Processed transcript	-	-	TSL:3
Rel1-202	ENSMUST00000127314.1	375	No protein	Processed transcript	-	-	TSL:3
Rel1-203	ENSMUST00000128967.1	494	No protein	Retained intron	-	-	TSL:2

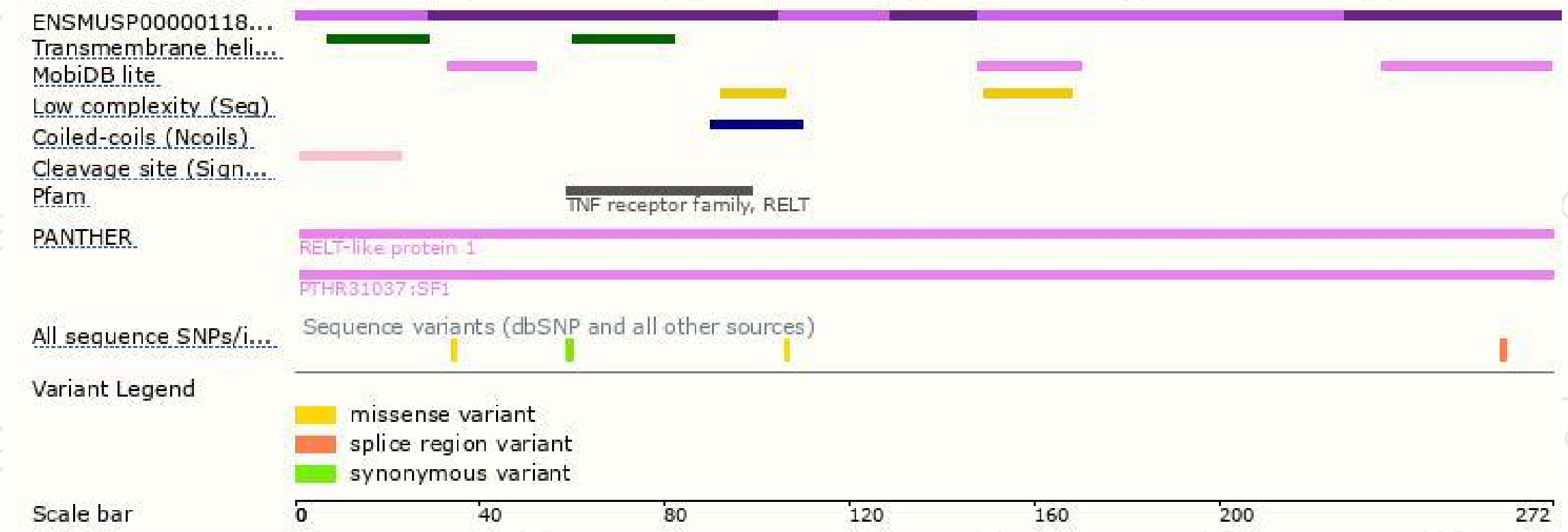
The strategy is based on the design of *Rel1-204* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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