

Copb1 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Copb1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Copb1* gene. The schematic diagram is as follows:



- The *Copb1* gene has 3 transcripts. According to the structure of *Copb1* gene, exon3-exon5 of *Copb1-201* (ENSMUST00000033012.8) transcript is recommended as the knockout region. The region contains 515bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Copb1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Copb1* gene is located on the Chr7. 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)

Copb1 coatomer protein complex, subunit beta 1 [Mus musculus (house mouse)]

Gene ID: 70349, updated on 7-Apr-2019

Summary



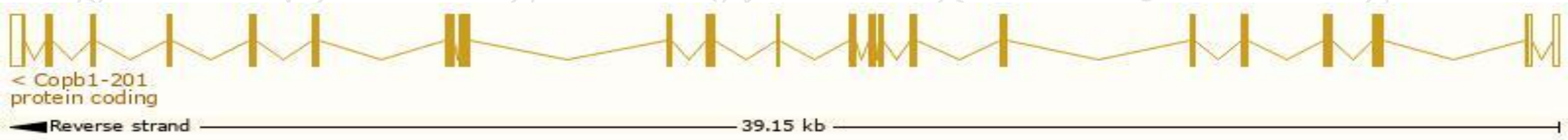
Official Symbol	Copb1 provided by MGI
Official Full Name	coatomer protein complex, subunit beta 1 provided by MGI
Primary source	MGI:MGI:1917599
See related	Ensembl:ENSMUSG00000030754
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2610019B04Rik
Expression	Ubiquitous expression in placenta adult (RPKM 44.6), limb E14.5 (RPKM 34.6) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

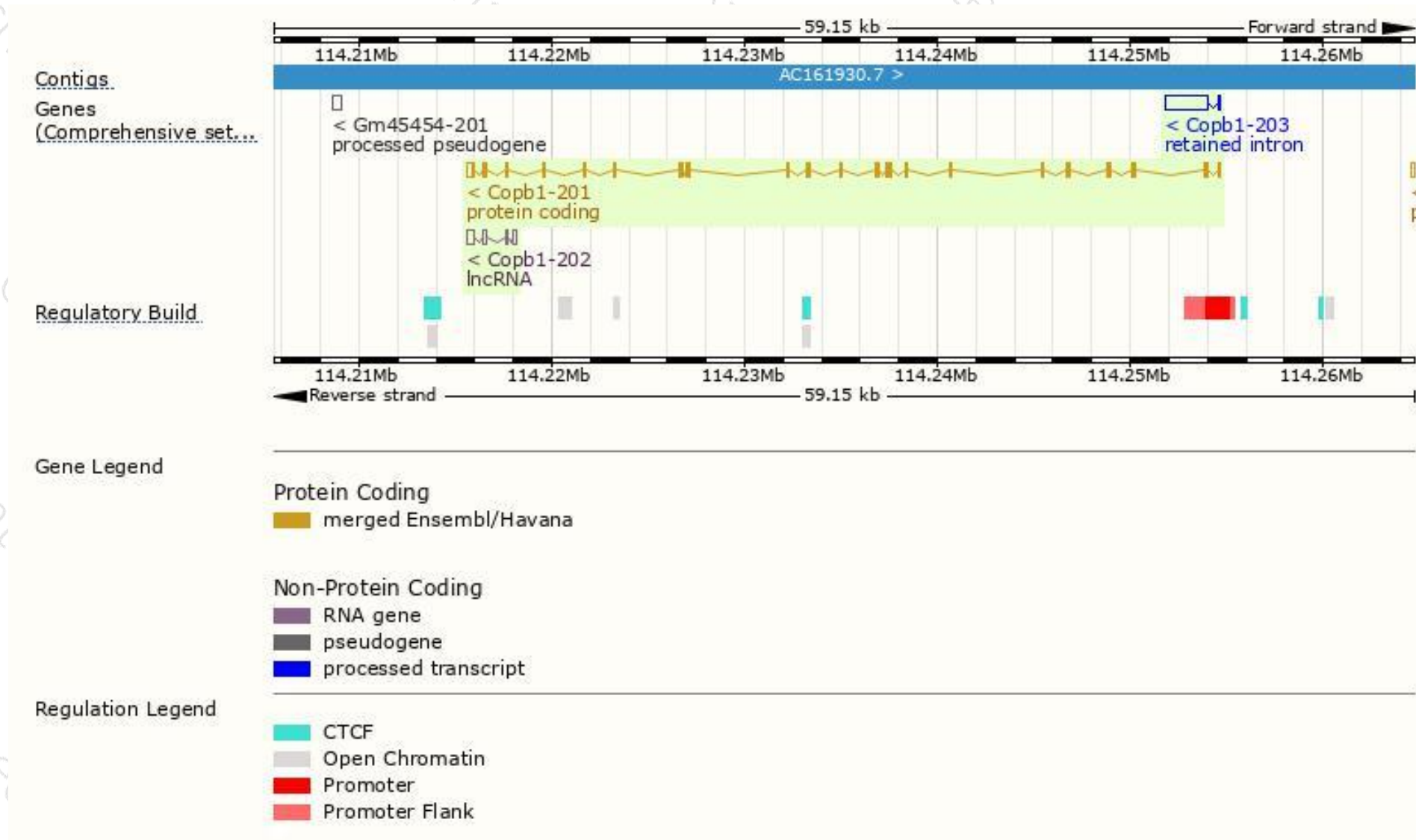
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Copb1-201	ENSMUST00000033012.8	3346	953aa	Protein coding	CCDS21759	Q9JIF7	TSL:1 GENCODE basic APPRIS P1
Copb1-203	ENSMUST00000211417.1	2302	No protein	Retained intron	-	-	TSL:1
Copb1-202	ENSMUST00000209394.1	760	No protein	lncRNA	-	-	TSL:3

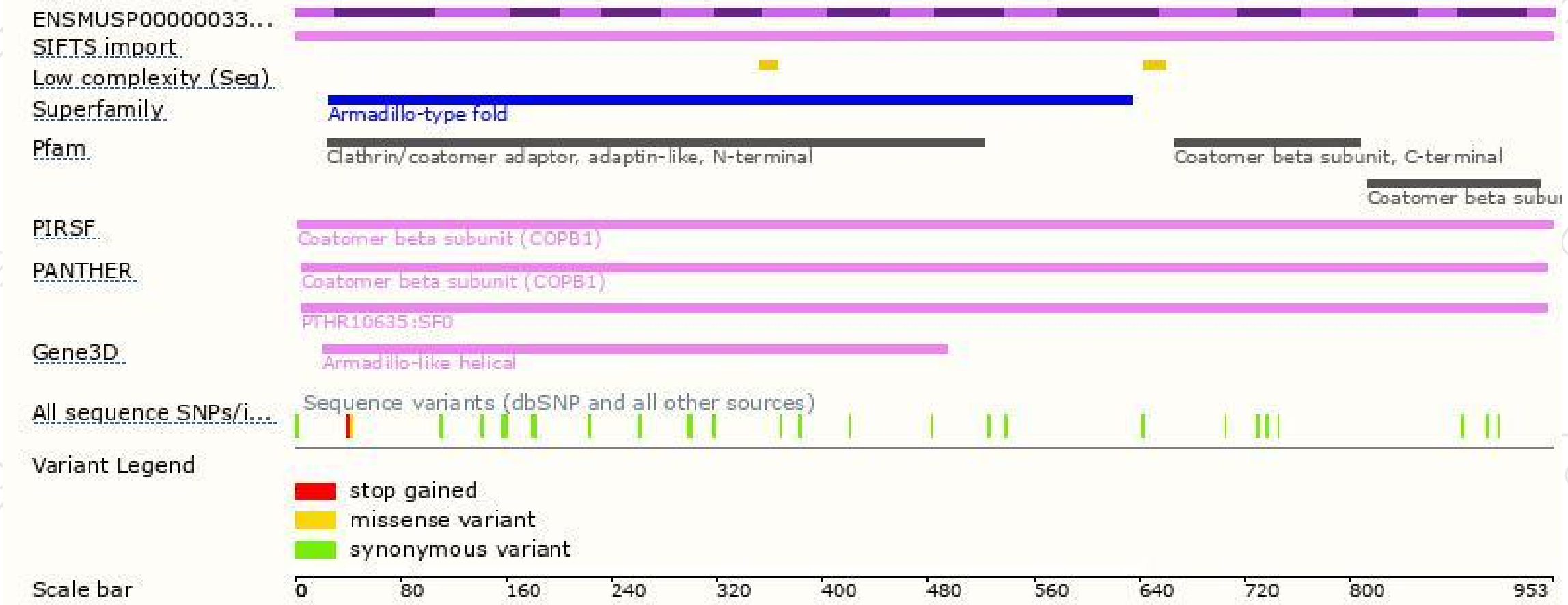
The strategy is based on the design of *Copb1-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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