

Eif3j1 Cas9-KO Strategy

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Project Overview

Project Name

Eif3j1

Project type

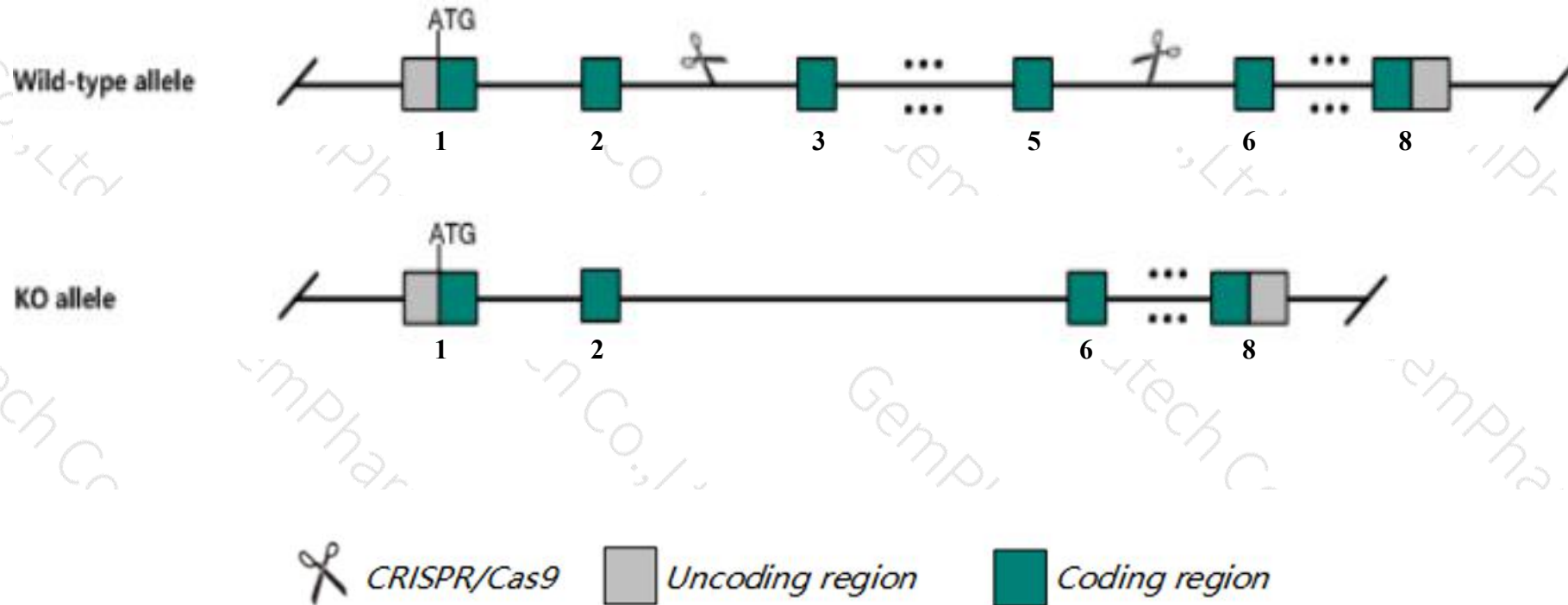
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Eif3j1* gene. The schematic diagram is as follows:



- The *Eif3j1* gene has 2 transcripts. According to the structure of *Eif3j1* gene, exon3-exon5 of *Eif3j1-201* (ENSMUST00000028668.7) transcript is recommended as the knockout region. The region contains 262bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Eif3j1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Eif3j1* gene is located on the Chr2. 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)

Eif3j1 eukaryotic translation initiation factor 3, subunit J1 [Mus musculus (house mouse)]

Gene ID: 78655, updated on 13-Mar-2020

Summary



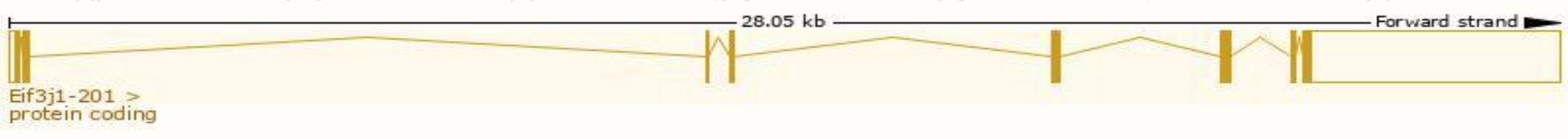
Official Symbol	Eif3j1 provided by MGI
Official Full Name	eukaryotic translation initiation factor 3, subunit J1 provided by MGI
Primary source	MGI:MGI:1925905
See related	Ensembl:ENSMUSG00000027236
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1810016I04Rik, 2700079K05Rik, AA409117, AA409446, C78575, Eif3j, Eif3j-1, Eif3j-2, Eif3j2, Eif3s1, eIF-3-alpha-A, eIF-3-alpha-B, eIF3-alpha, eIF3-p35, eIF3j-A, eIF3j-B
Expression	Broad expression in CNS E11.5 (RPKM 29.5), liver E14 (RPKM 27.9) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

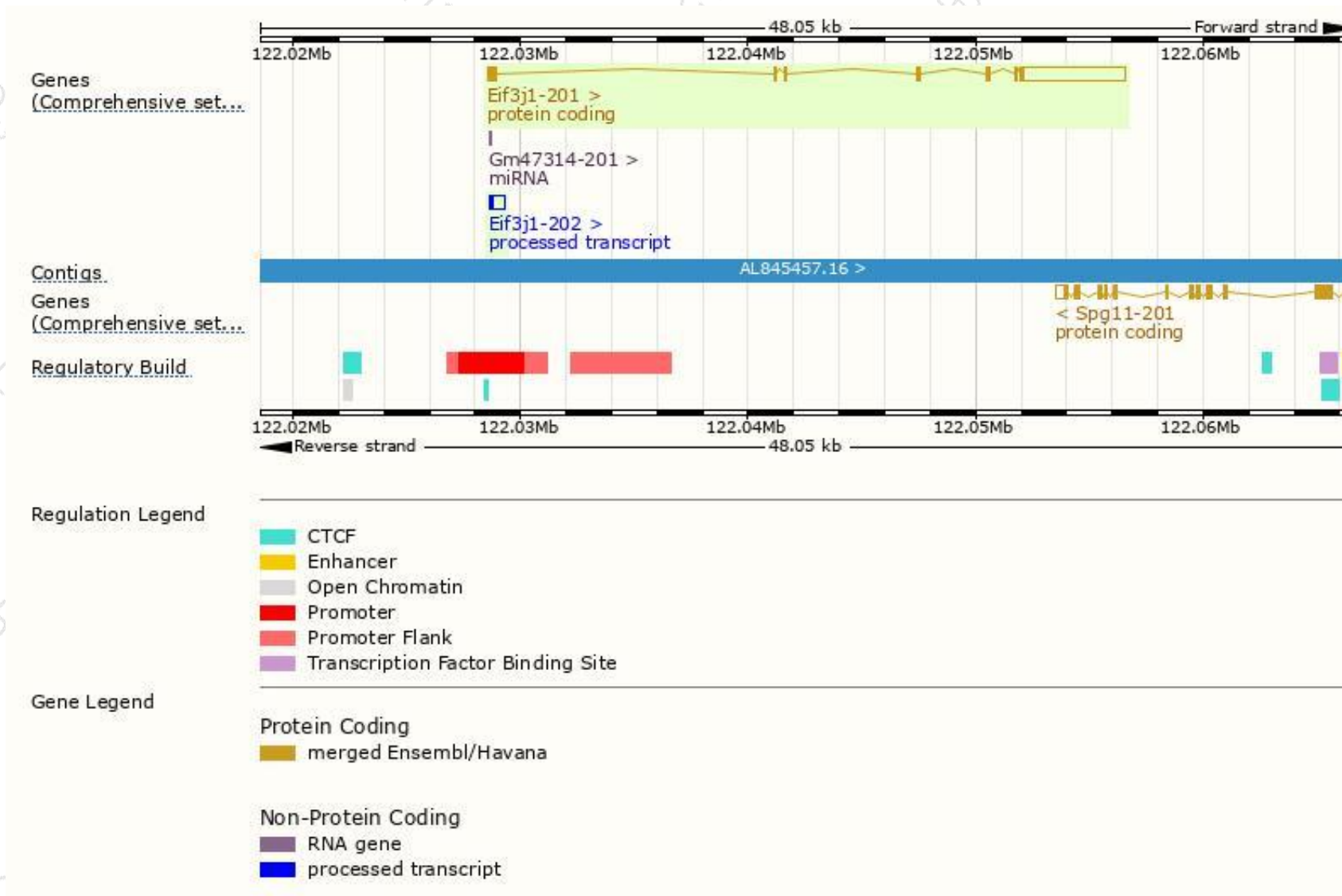
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Eif3j1-201	ENSMUST00000028668.7	5413	261aa	Protein coding	CCDS38219	Q3UGC7	TSL:1 GENCODE basic APPRIS P1
Eif3j1-202	ENSMUST00000142747.1	594	No protein	Processed transcript	-	-	TSL:2

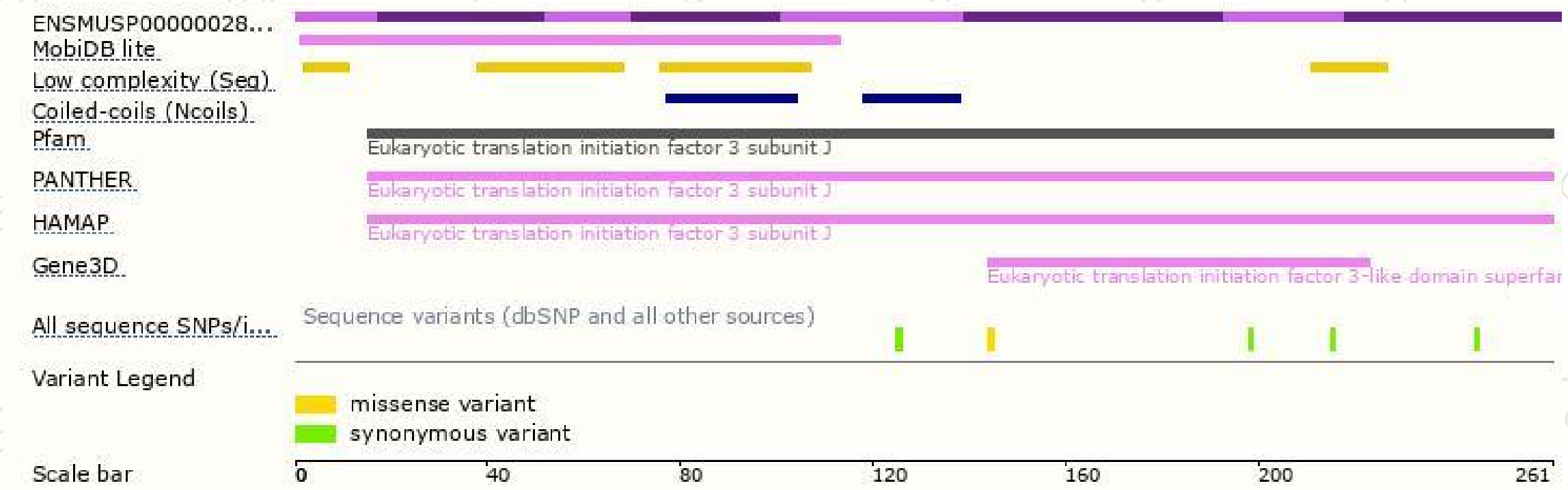
The strategy is based on the design of *Eif3j1-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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