

Eif3l Cas9-CKO Strategy

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Project Overview

Project Name

Eif3l

Project type

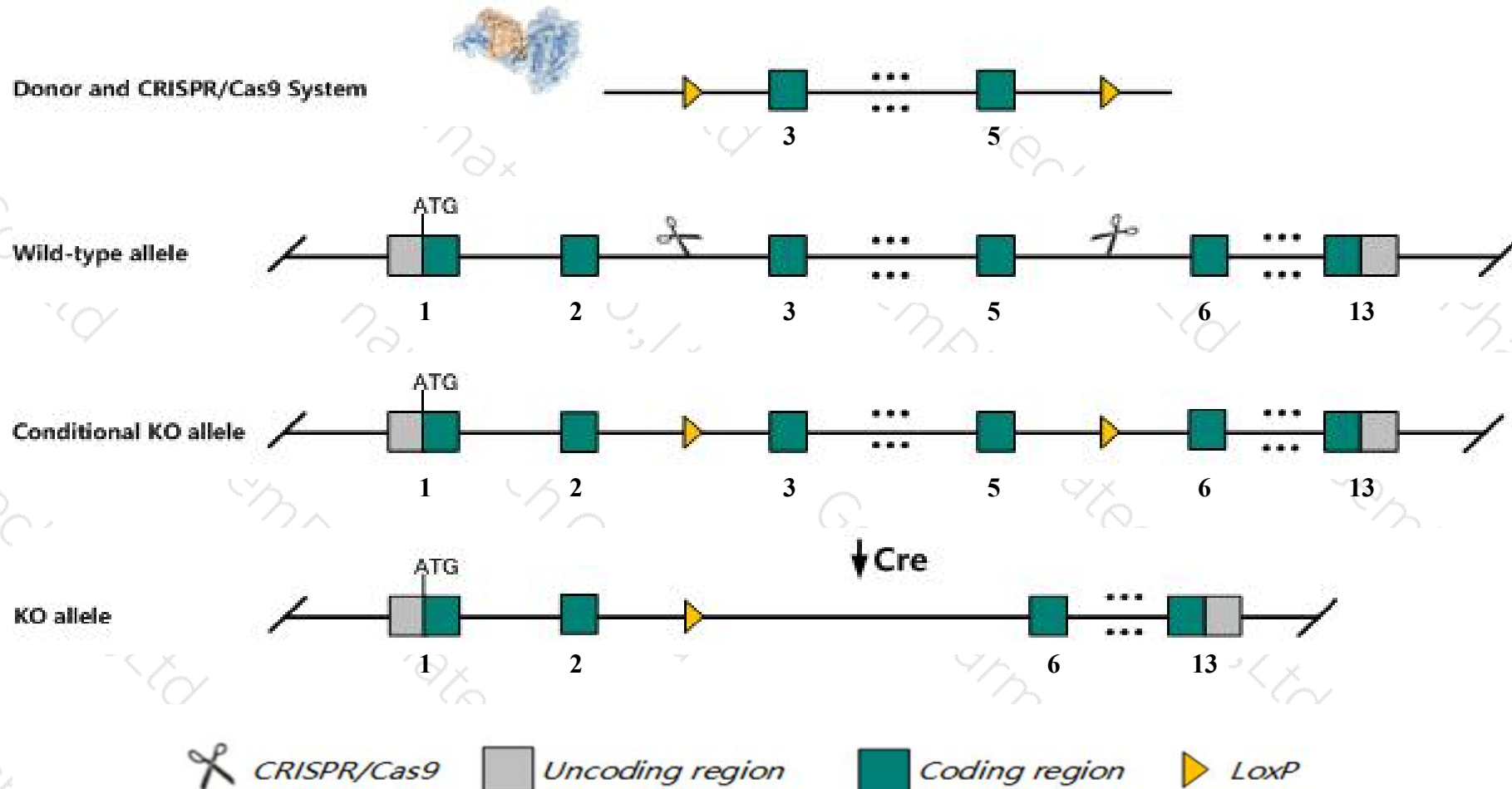
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Eif3l* gene. The schematic diagram is as follows:



Technical routes

- The *Eif3l* gene has 4 transcripts. According to the structure of *Eif3l* gene, exon3-exon5 of *Eif3l-201* (ENSMUST00000040518.5) transcript is recommended as the knockout region. The region contains 353bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Eif3l* gene. The brief process is as follows: CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- The *Eif3l* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Eif3l eukaryotic translation initiation factor 3, subunit L [Mus musculus (house mouse)]

Gene ID: 223691, updated on 13-Mar-2020

Summary



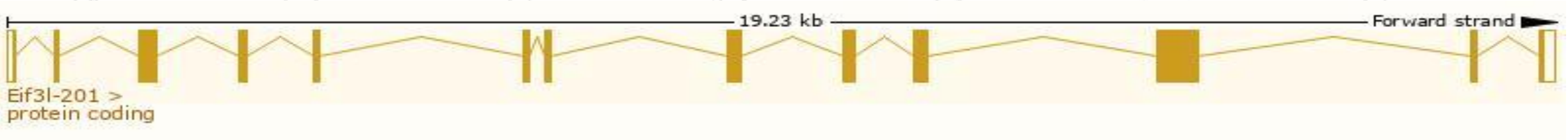
Official Symbol	Eif3l provided by MGI
Official Full Name	eukaryotic translation initiation factor 3, subunit L provided by MGI
Primary source	MGI:MGI:2386251
See related	Ensembl:ENSMUSG00000033047
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	0610011H21Rik, D15N1e, Eif3eip, Eif3ip, Eif3s6ip, HSP-66Y, PAF67
Expression	Ubiquitous expression in placenta adult (RPKM 103.9), ovary adult (RPKM 96.4) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

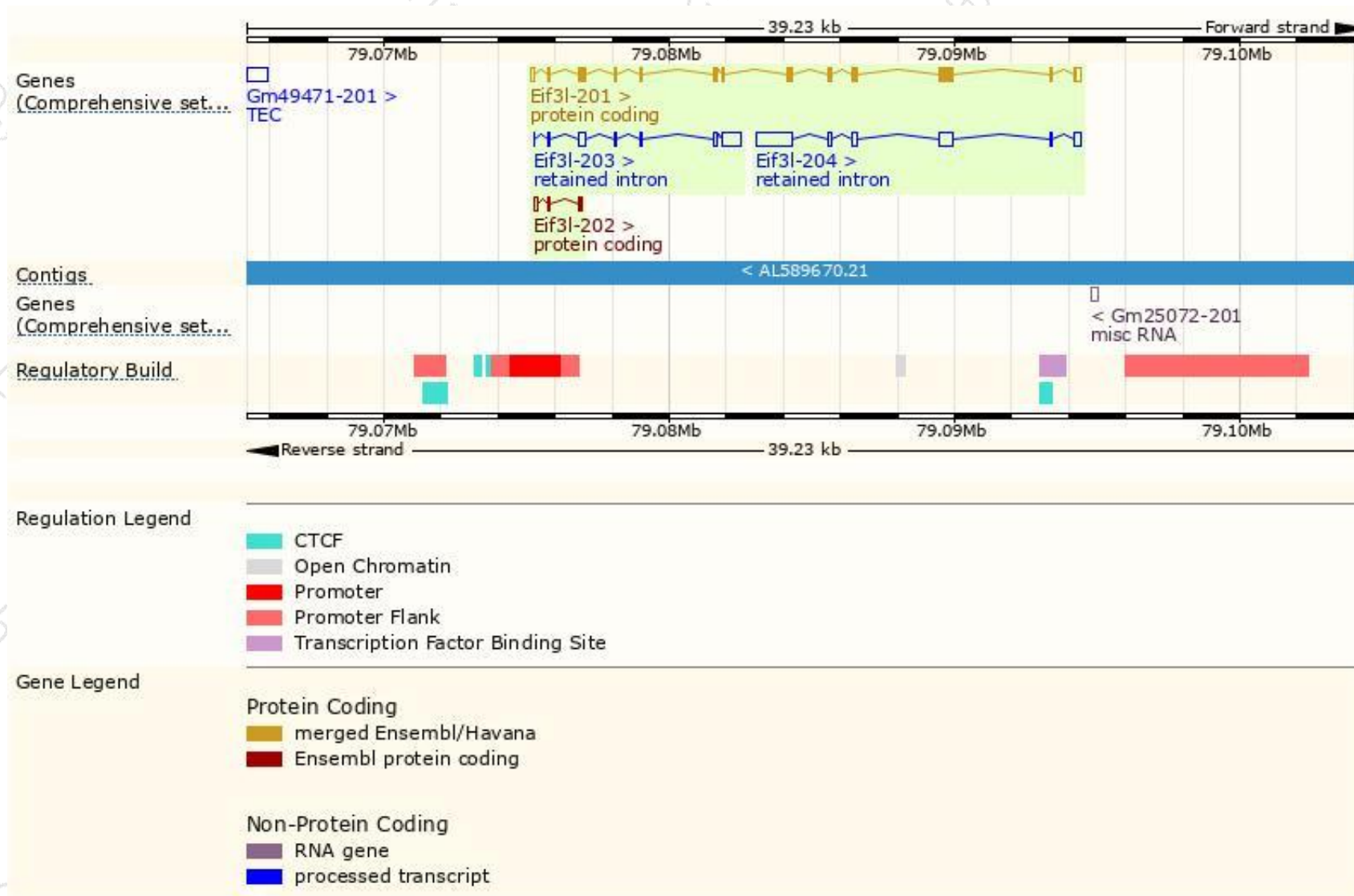
The gene has 4 transcripts,all transcripts are shown below:

Name ▲	Transcript ID ▲	bp ▲	Protein ▲	Biotype ▲	CCDS ▲	UniProt ▲	Flags ▲
Eif3l-201	ENSMUST00000040518.5	1942	564aa	Protein coding	CCDS27632	Q8QZY1	TSL:1 GENCODE basic APPRIS P1
Eif3l-202	ENSMUST000000229310.1	274	44aa	Protein coding	-	A0A2R8VH97	CDS 3' incomplete
Eif3l-203	ENSMUST000000229338.1	1157	No protein	Retained intron	-	-	-
Eif3l-204	ENSMUST000000230032.1	2345	No protein	Retained intron	-	-	-

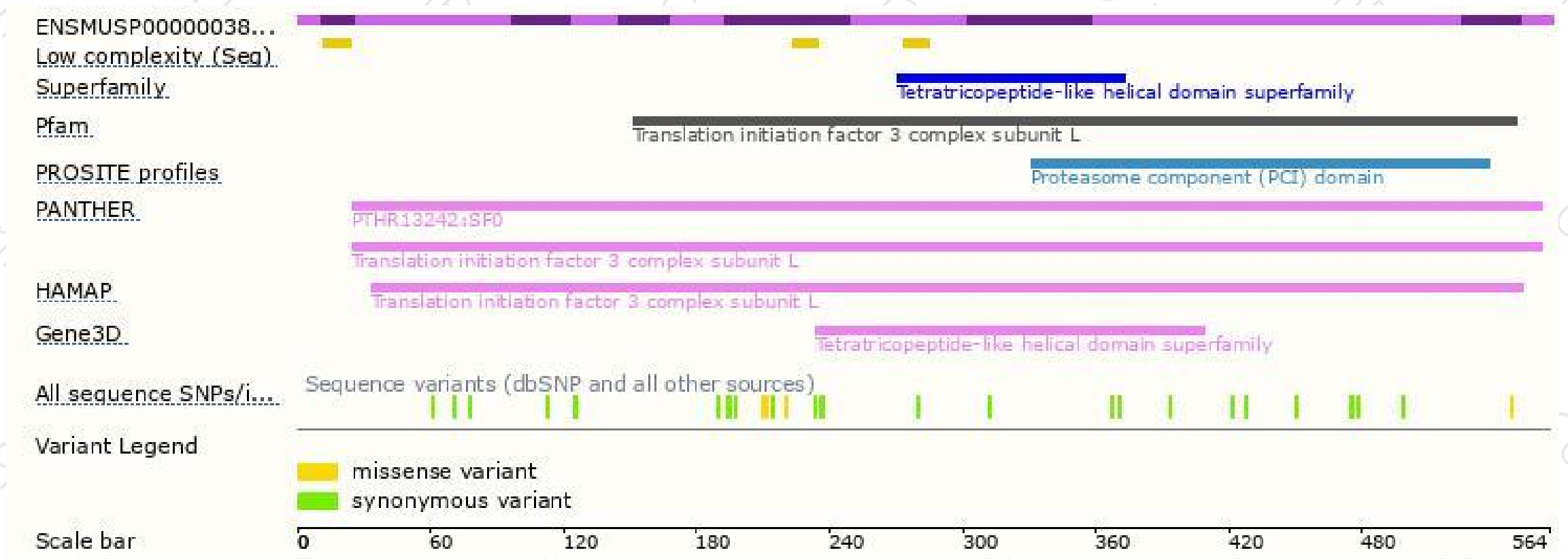
The strategy is based on the design of *Eif3l-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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