

Slc25a18 Cas9-CKO Strategy

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Project Overview

Project Name

Slc25a18

Project type

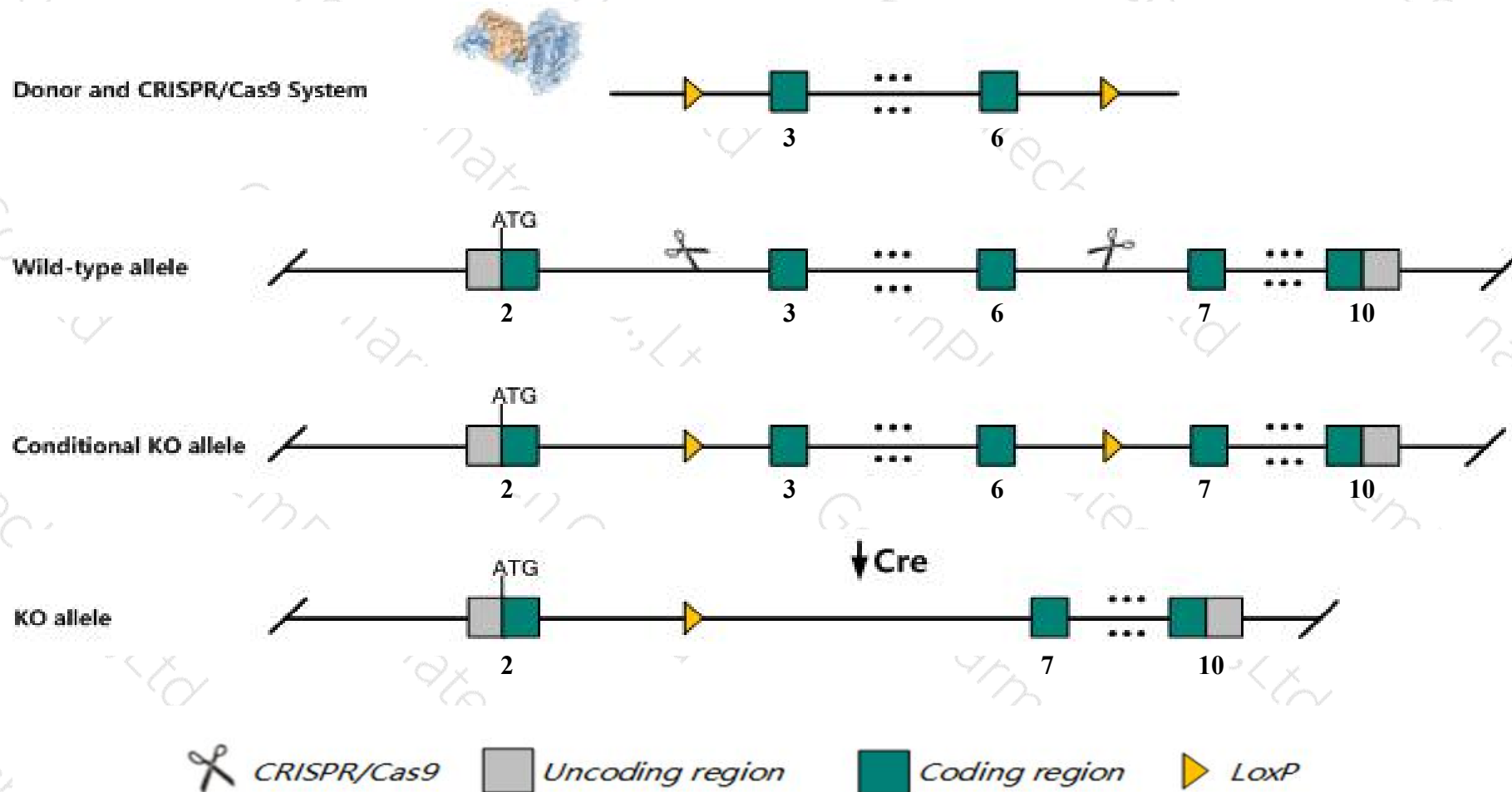
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc25a18* gene. The schematic diagram is as follows:



Technical routes

- The *Slc25a18* gene has 1 transcript. According to the structure of *Slc25a18* gene, exon3-exon6 of *Slc25a18-201* (ENSMUST00000112682.3) transcript is recommended as the knockout region. The region contains 389bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc25a18* 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 *Slc25a18* gene is located on the Chr6. 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)



Slc25a18 solute carrier family 25 (mitochondrial carrier), member 18 [*Mus musculus* (house mouse)]

Gene ID: 71803, updated on 12-Aug-2019

Summary

- Official Symbol

Slc25a18 provided by MGI
- Official Full Name

solute carrier family 25 (mitochondrial carrier), member 18 provided by MGI
- Primary source

[MGI:MGI:1919053](#)
- See related

[Ensembl:ENSMUSG00000004902](#)
- 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

AW125787; 1500015I14Rik
- Expression

Biased expression in cortex adult (RPKM 19.5), frontal lobe adult (RPKM 19.4) and 6 other tissues [See more](#)
- Orthologs

[human](#) [all](#)

Genomic context

Location: 6 F1; 6 57.0 cM

See Slc25a18 in [Genome Data Viewer](#)

Exon count: 11

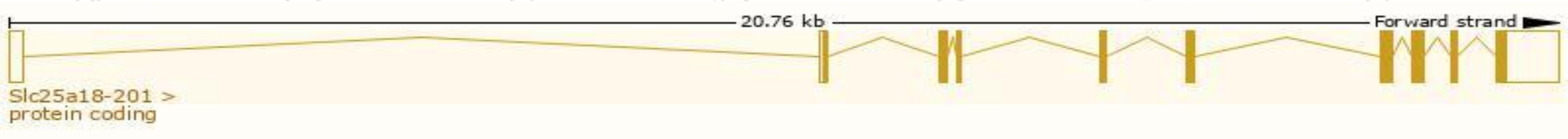
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	6	NC_000072.6 (120773504..120794169)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	6	NC_000072.5 (120723786..120744000)

Transcript information (Ensembl)

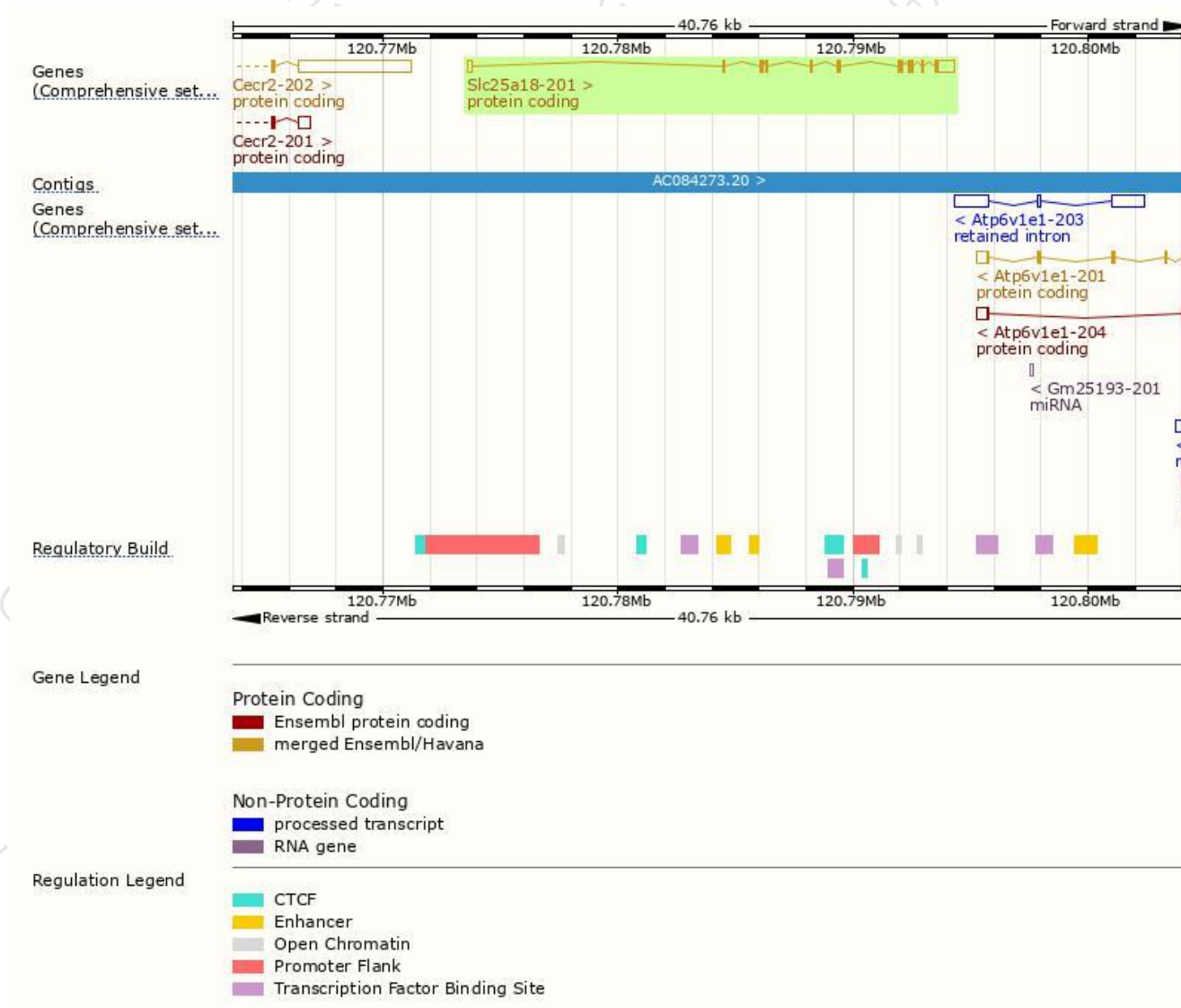
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc25a18-201	ENSMUST00000112682.3	1911	320aa	Protein coding	CCDS39614	Q9DB41	TSL:1 GENCODE basic APPRIS P1

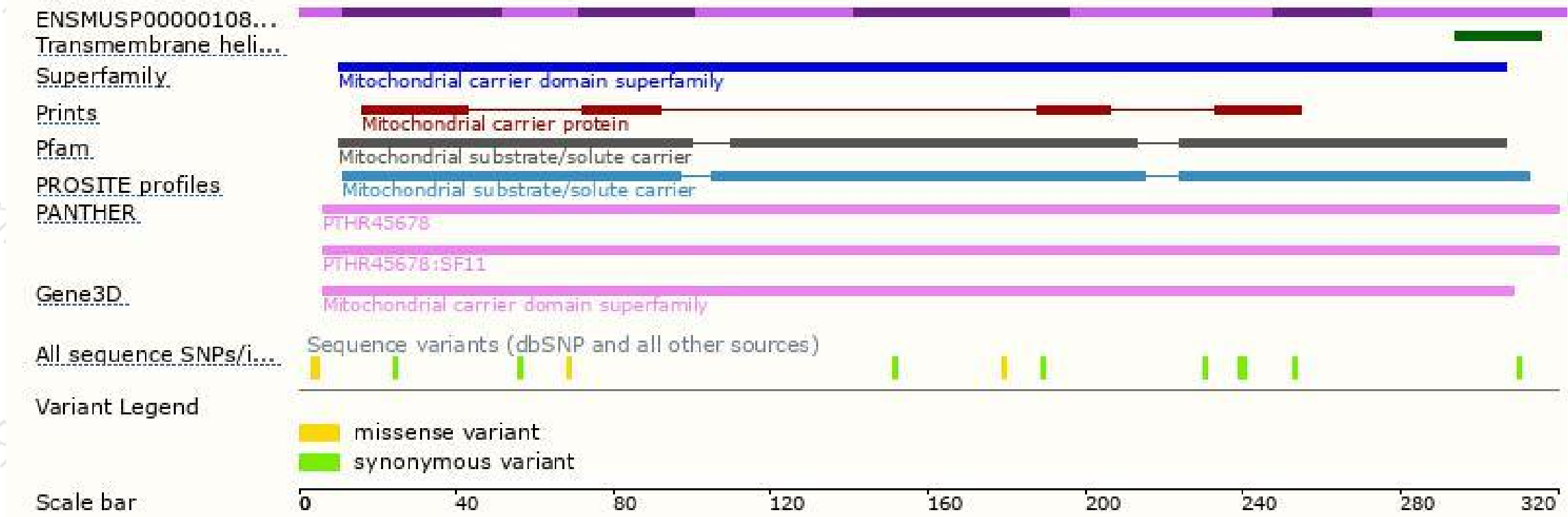
The strategy is based on the design of *Slc25a18-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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