

Slc25a48 Cas9-CKO Strategy

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Design Date: 2021-7-9

Project Overview

Project Name

Slc25a48

Project type

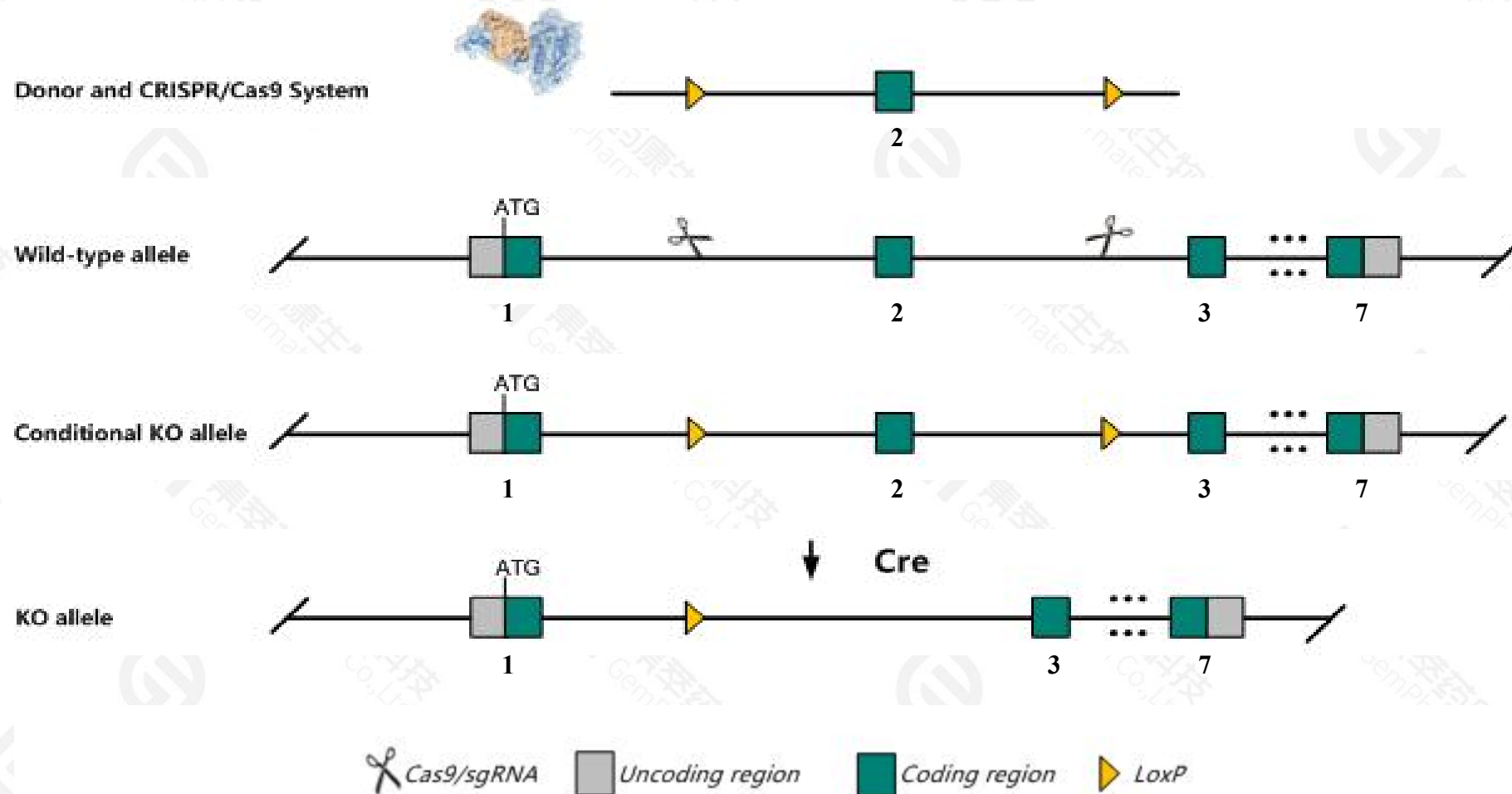
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc25a48* gene. The schematic diagram is as follows:



Technical routes

- The *Slc25a48* gene has 1 transcript. According to the structure of *Slc25a48* gene, exon2 of *Slc25a48*-201(ENSMUST00000021971.6) transcript is recommended as the knockout region. The region contains 44bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc25a48* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor was constructed. Cas9, sgRNA 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 was 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.

- The *Slc25a48* gene is located on the Chr13. 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)

Slc25a48 solute carrier family 25, member 48 [Mus musculus (house mouse)]

Gene ID: 328258, updated on 17-Dec-2020

Summary



Official Symbol Slc25a48 provided by [MGI](#)

Official Full Name solute carrier family 25, member 48 provided by [MGI](#)

Primary source [MGI:MGI:2145373](#)

See related [Ensembl:ENSMUSG00000021509](#)

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 AU042651, E230025K15

Expression Biased expression in ovary adult (RPKM 14.5), subcutaneous fat pad adult (RPKM 6.6) and 7 other tissues [See more](#)

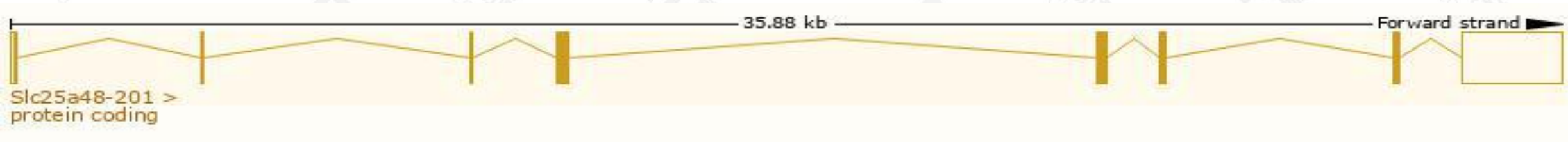
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

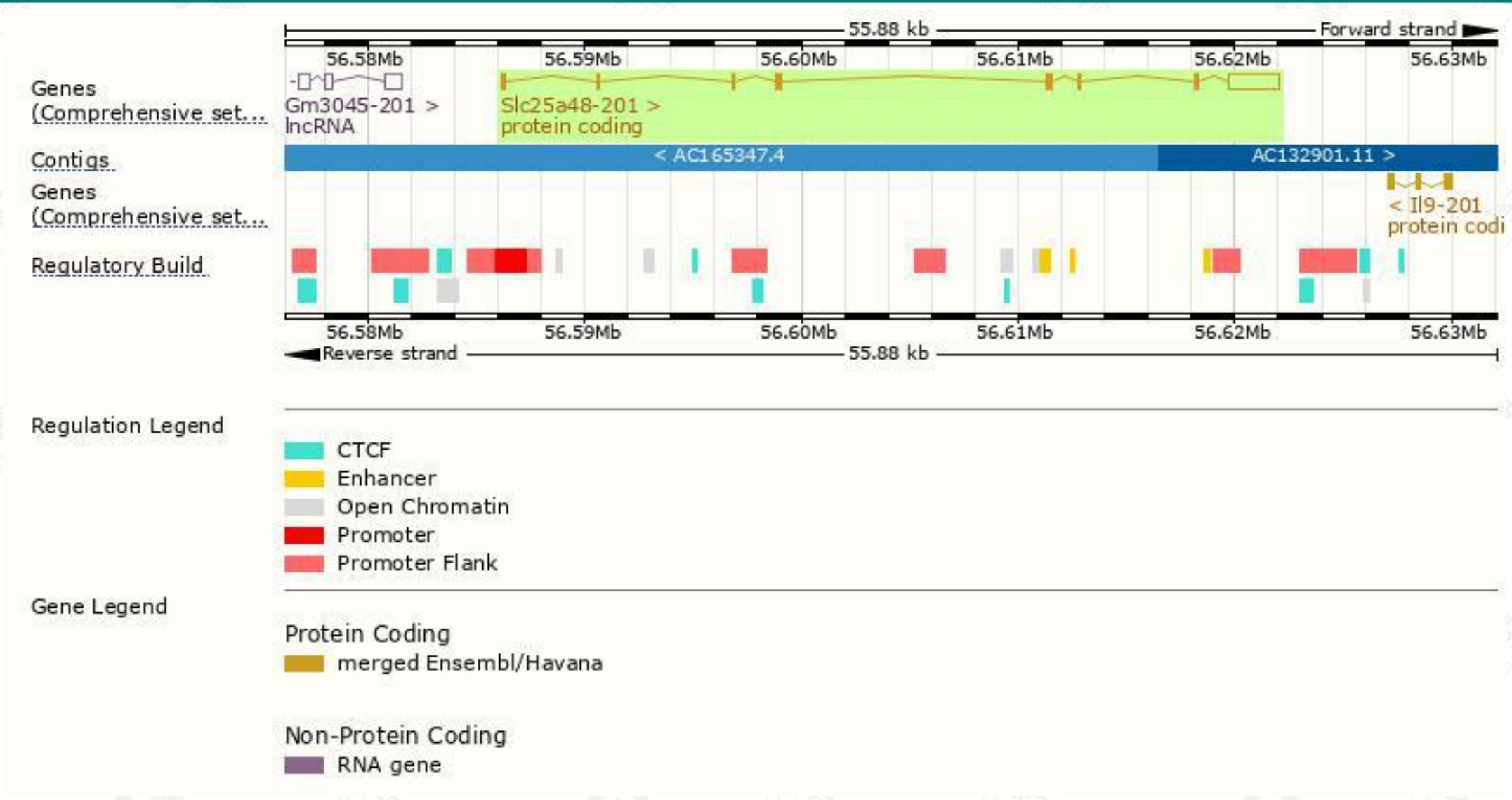
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc25a48-201	ENSMUST00000021971.6	3374	306aa	Protein coding	CCDS26561		TSL:1 , GENCODE basic , APPRIS P1 ,

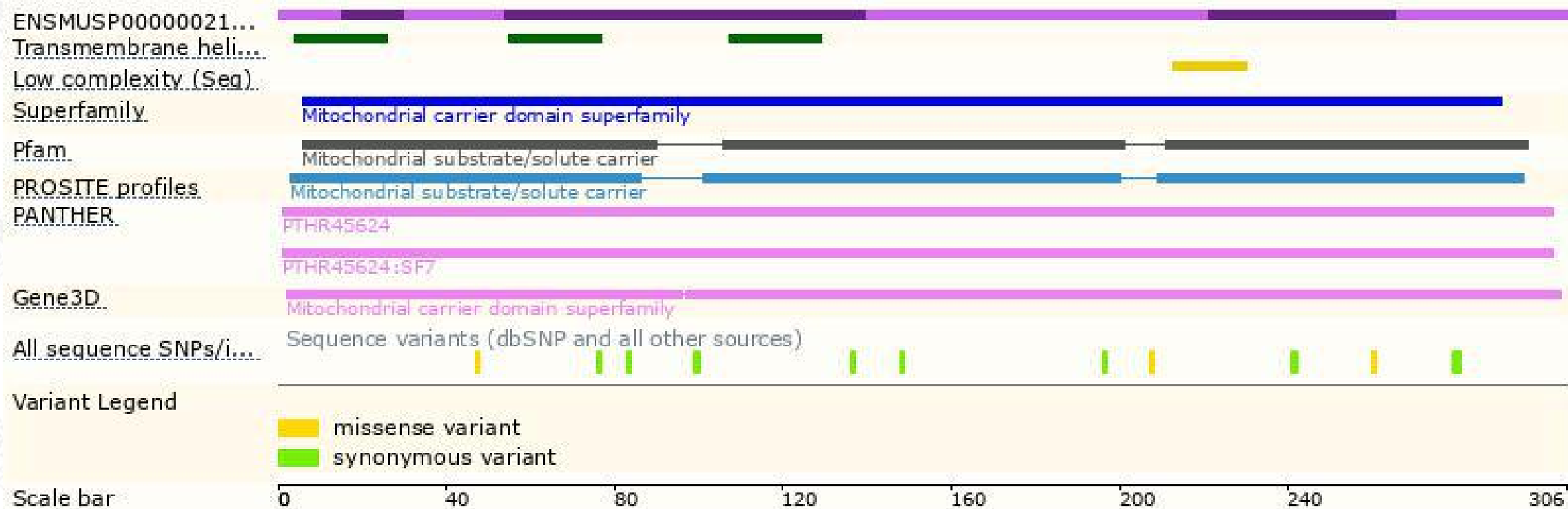
The strategy is based on the design of *Slc25a48-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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