Slc25a4 Cas9-KO Strategy

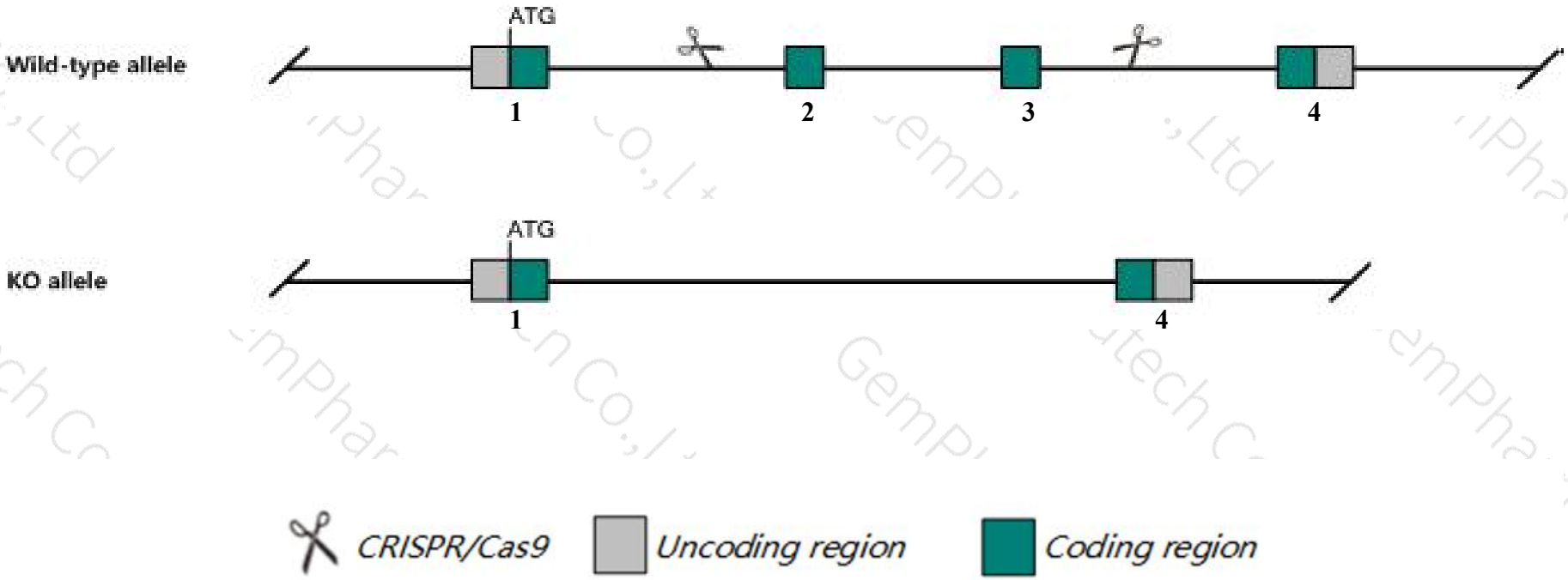
Designer: Lixin Lv

Project Overview

Project Name	***Slc25a4***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc25a4* gene. The schematic diagram is as follows:



Technical routes

➢ The *Slc25a4* gene has 2 transcripts. According to the structure of *Slc25a4* gene, exon2-exon3 of *Slc25a4-201* (ENSMUST00000034049.4) transcript is recommended as the knockout region.The region contains 628bp coding sequence. Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Slc25a4* gene. The brief process is as follows: CRISPR/Cas9 syste

Notice

➢ According to the existing MGI data,Homozygous null mice exhibit a defect in mitochondrial energy metabolism and develop mitochondrial myopathy and hypertrophic cardiomyopathy, metabolic acidosis, and a severe exercise intolerance.

➢ The *Slc25a4* gene is located on the Chr8. 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）

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Slc25a4 solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4 [Mus musculus (house mouse)]

Gene ID: 11739, updated on 7-Apr-2019

Summary

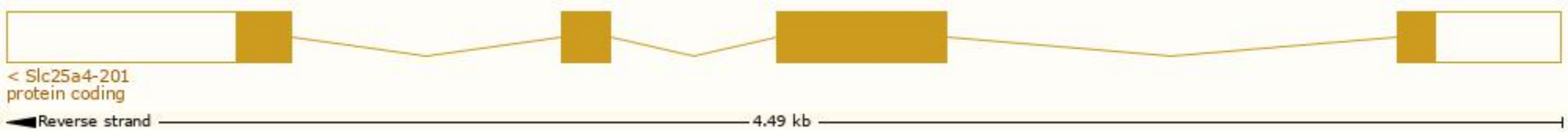
Official Symbol	Slc25a4 provided by MGI
Official Full Name	solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4 provided by MGI
Primary source	MGI:MGI:1353495
See related	Ensembl:ENSMUSG00000031633
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AU019225, Ant1, mANC1
Expression	Broad expression in heart adult (RPKM 2616.6), cerebellum adult (RPKM 363.2) and 16 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

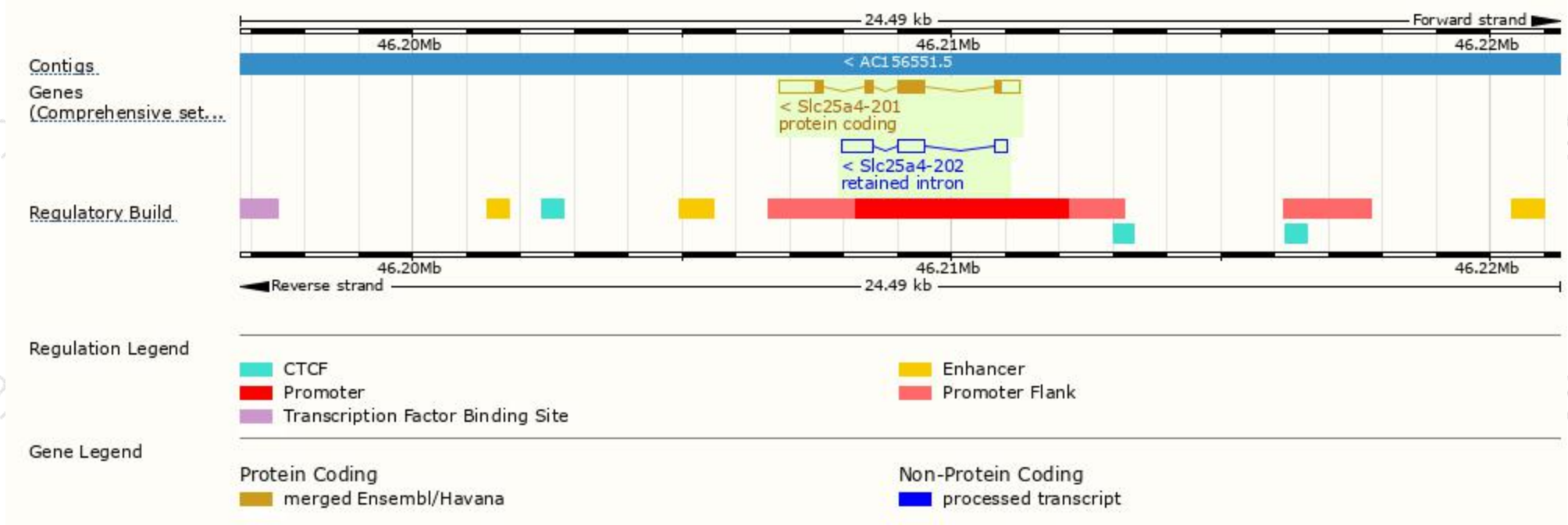
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc25a4-201	ENSMUST00000034049.4	1925	298aa	Protein coding	CCDS40333	P48962	TSL:1 GENCODE basic APPRIS P1
Slc25a4-202	ENSMUST00000155986.1	1275	No protein	Retained intron	-	-	TSL:1

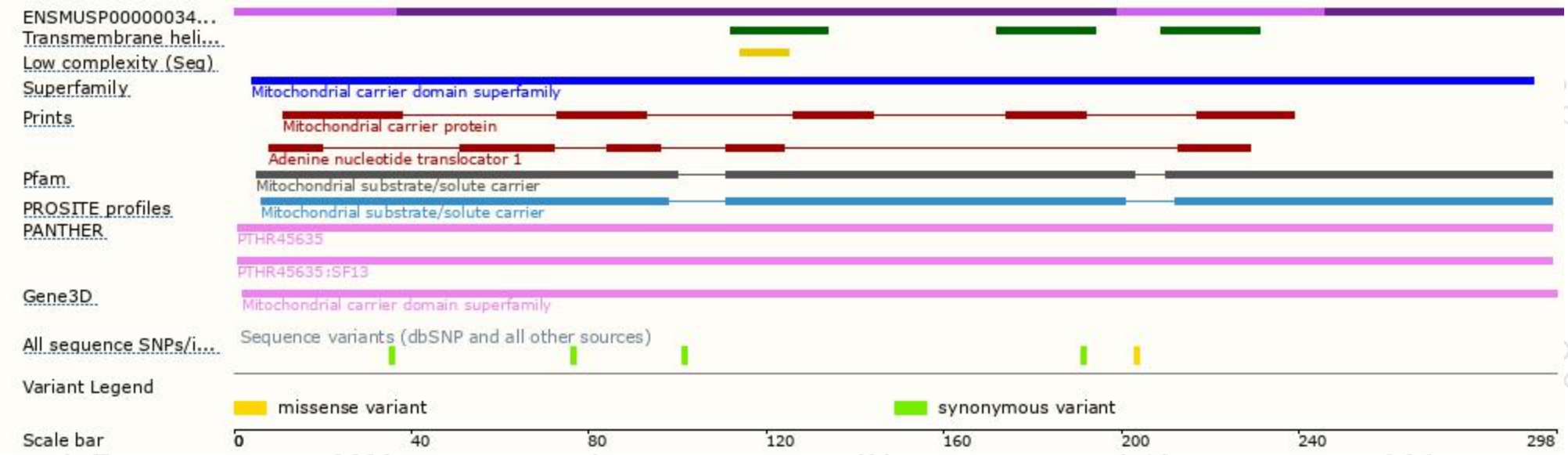
The strategy is based on the design of *Slc25a4-201* transcript,The transcription is shown below



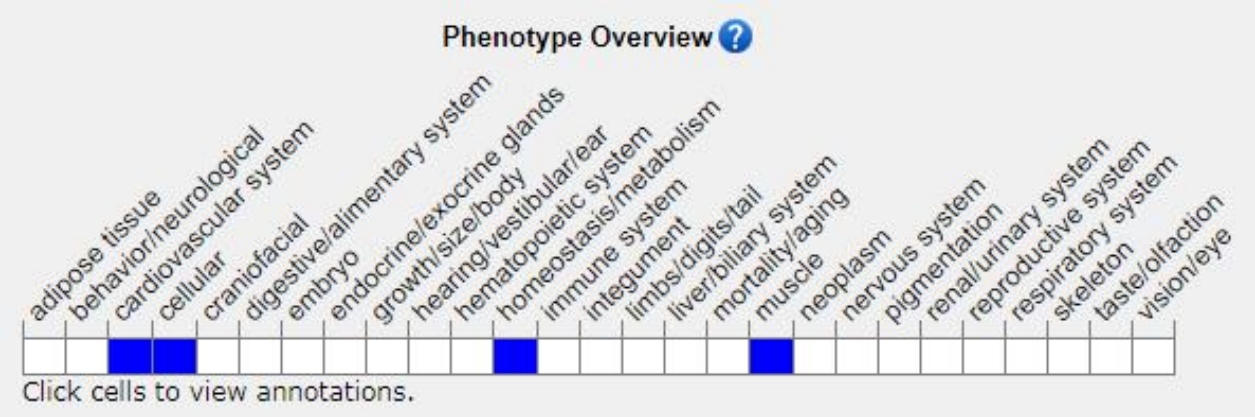
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

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

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