

# Slc25a11 Cas9-KO Strategy

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# Overview

## Target Gene Name

- Slc25a11

## Project Type

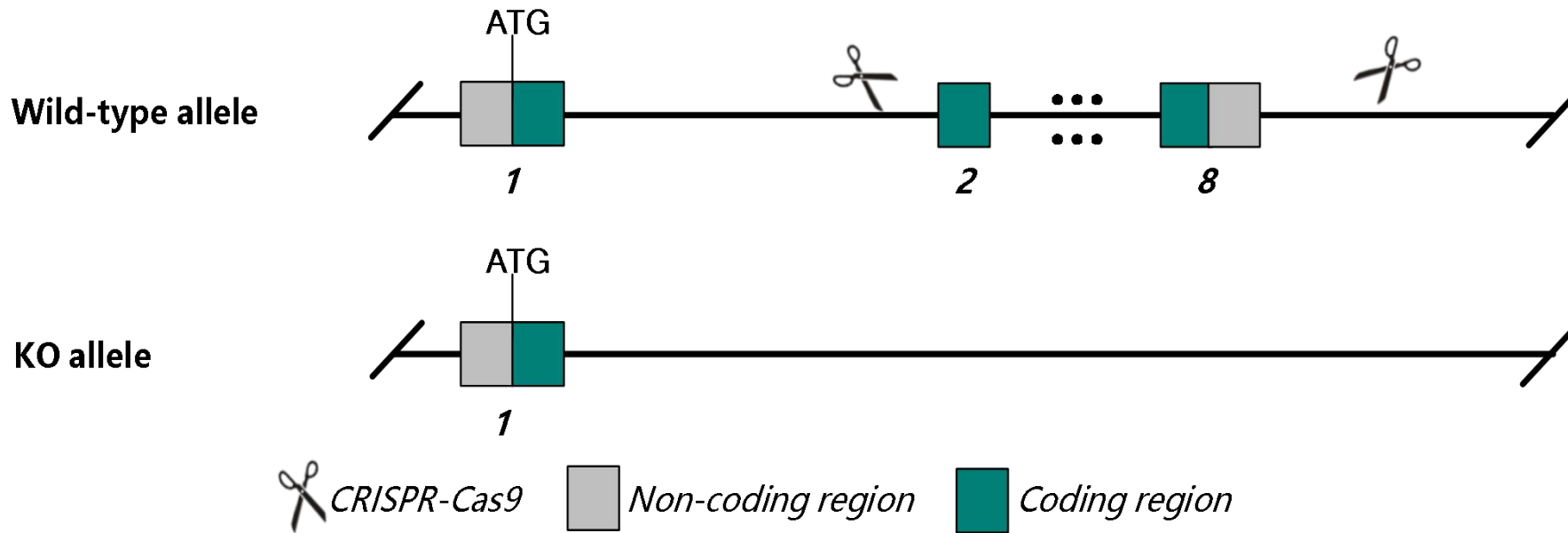
- Cas9-KO

## Genetic Background

- C57BL/6JGpt



# Strain Strategy



Schematic representation of CRISPR-Cas9 engineering used to edit the *Slc25a11* gene.



# Technical Information

- The *Slc25a11* gene has 5 transcripts. According to the structure of *Slc25a11* gene, exon2-8 of *Slc25a11*-201 (ENSMUST00000014750.15) transcript is recommended as the knockout region. The region contains most of coding sequences. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Slc25a11* gene. The brief process is as follows: gRNAs were transcribed in vitro. Cas9 and gRNAs 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 on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.



# Gene Information

**Slc25a11** solute carrier family 25 (mitochondrial carrier oxoglutarate carrier), member 11 [ *Mus musculus* (house mouse) ]

Gene ID: 67863, updated on 14-Aug-2022

[Download Datasets](#)

## Summary

<b>Official Symbol</b>	Slc25a11 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	solute carrier family 25 (mitochondrial carrier oxoglutarate carrier), member 11 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1915113</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000014606</a> <a href="#">AllianceGenome:MGI:1915113</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	2oxoc; 2310022P18Rik
<b>Summary</b>	Predicted to enable antiporter activity; dicarboxylic acid transmembrane transporter activity; and sulfur compound transmembrane transporter activity. Predicted to be involved in anion transport. Located in mitochondrial inner membrane. Is expressed in embryo. Human ortholog(s) of this gene implicated in paraganglioma. Orthologous to human SLC25A11 (solute carrier family 25 member 11). [provided by Alliance of Genome Resources, Apr 2022]
<b>Expression</b>	Ubiquitous expression in heart adult (RPKM 208.5), kidney adult (RPKM 85.6) and 27 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>
<b>NEW</b>	Try the new <a href="#">Gene table</a> Try the new <a href="#">Transcript table</a>

## Genomic context

**Location:** 11 B3; 11 43.21 cM

**Exon count:** 9

See Slc25a11 in [Genome Data Viewer](#)

Source: <https://www.ncbi.nlm.nih.gov/>

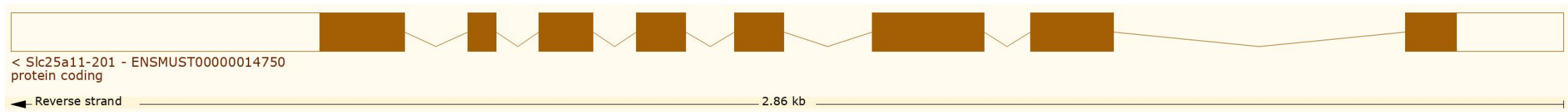


# Transcript Information

The gene has 5 transcripts, all transcripts are shown below:

Transcript ID	Name	bp	Protein	Biotype	CCDS	UniProt Match	Flags
<a href="#">ENSMUST00000014750.15</a>	Slc25a11-201	1711	<a href="#">314aa</a>	Protein coding	<a href="#">CCDS24958</a>	<a href="#">Q5SX53</a> <a href="#">Q9CR62</a>	Ensembl Canonical Gencode basic APPRIS P1 TSL:1
<a href="#">ENSMUST00000139638.2</a>	Slc25a11-204	770	<a href="#">193aa</a>	Protein coding		<a href="#">Q5SX46</a>	TSL:3 CDS 3' incomplete
<a href="#">ENSMUST00000136383.2</a>	Slc25a11-203	416	<a href="#">76aa</a>	Protein coding		<a href="#">Q5SX48</a>	TSL:3 CDS 3' incomplete
<a href="#">ENSMUST00000134804.2</a>	Slc25a11-202	768	No protein	Retained intron		-	TSL:2
<a href="#">ENSMUST00000157076.2</a>	Slc25a11-205	531	No protein	Retained intron		-	TSL:1

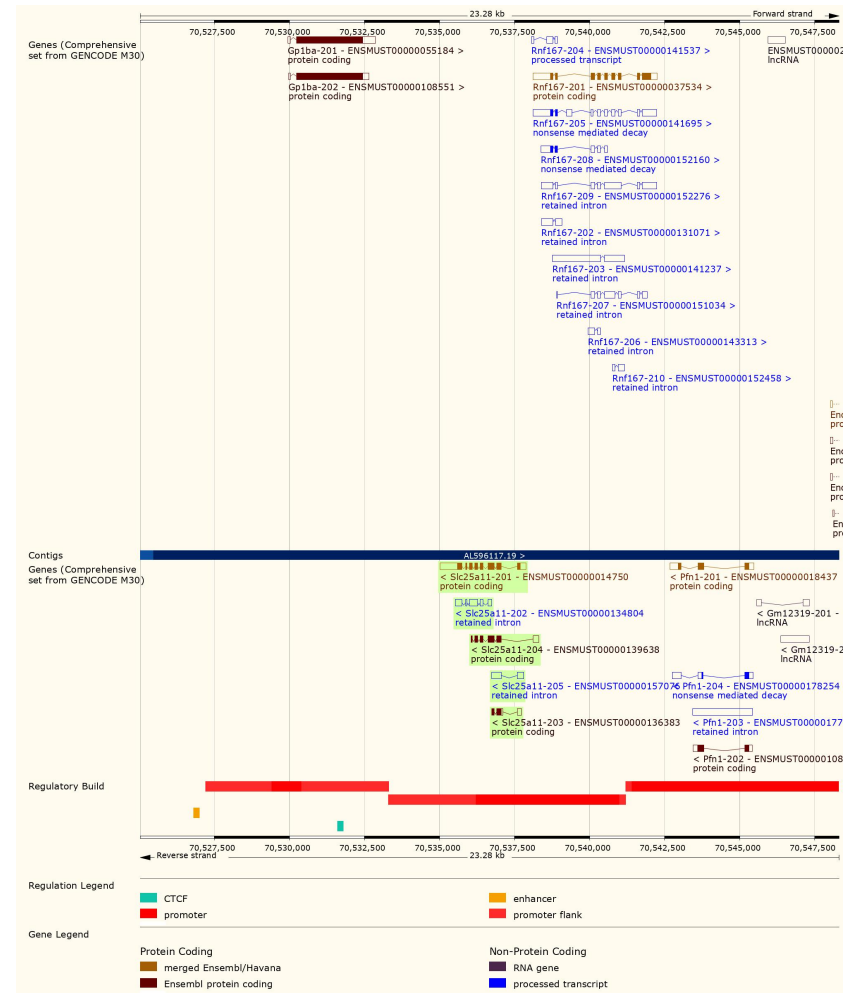
The strategy is based on the design of *Slc25a11-201* transcript, the transcription is shown below:



Source: <https://www.ensembl.org>

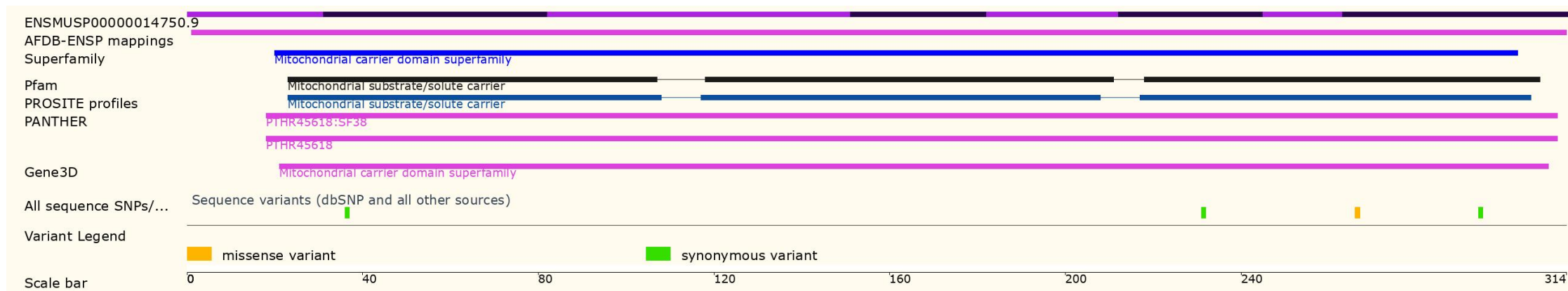


# Genomic Information





# Protein Information





# Mouse Phenotype Information (MGI)

- Homozygous knockout is embryonic lethal. Heterozygous KO reduces the tumor incidence of Kras activation mutation-induced lung tumors.



# Important Information

- According to MGI, homozygous knockout is embryonic lethal. Heterozygous KO reduces the tumor incidence of Kras activation mutation-induced lung tumors.
- The effect of *Rnf167*, *Gp1ba* and *Slc25a11*-203, *Slc25a11*-204 gene is unknown.
- *Slc25a11* is located on Chr11. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risks of the mutation on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.