

# Tomm22 Cas9-KO Strategy

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# Overview

## Target Gene Name

- Tomm22

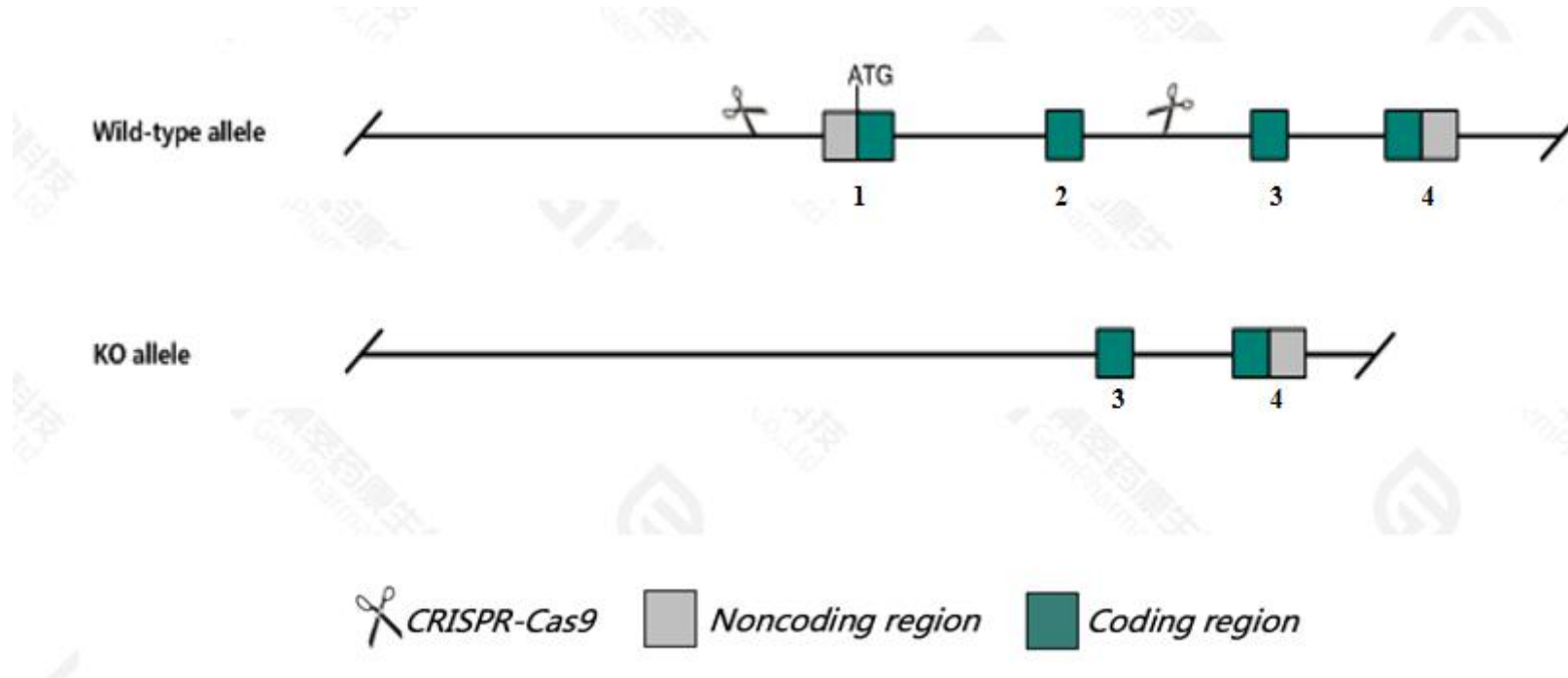
## Project Type

- Cas9-KO

## Genetic Background

- C57BL/6JGpt

# Strain Strategy



Schematic representation of CRISPR-Cas9 engineering used to edit the *Tom22* gene.

# Technical Information

- The *Tomm22* gene has 3 transcripts. According to the structure of *Tomm22* gene, exon1-exon2 of *Tomm22-201*(ENSMUST00000023062.5) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Tomm22* 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

## Tomm22 translocase of outer mitochondrial membrane 22 [Mus musculus (house mouse)]

Gene ID: 223696, updated on 24-Apr-2022

### Summary



<b>Official Symbol</b>	Tomm22 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	translocase of outer mitochondrial membrane 22 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2450248</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000022427</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	PROVISIONAL
<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>	2310047D01, Tom22
<b>Expression</b>	Ubiquitous expression in testis adult (RPKM 96.9), placenta adult (RPKM 50.2) and 28 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

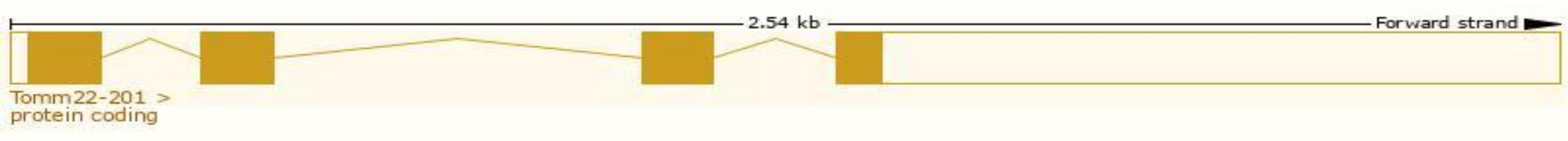
Source: <https://www.ncbi.nlm.nih.gov/>

# Transcript Information

The gene has 3 transcripts, all transcripts are shown below:

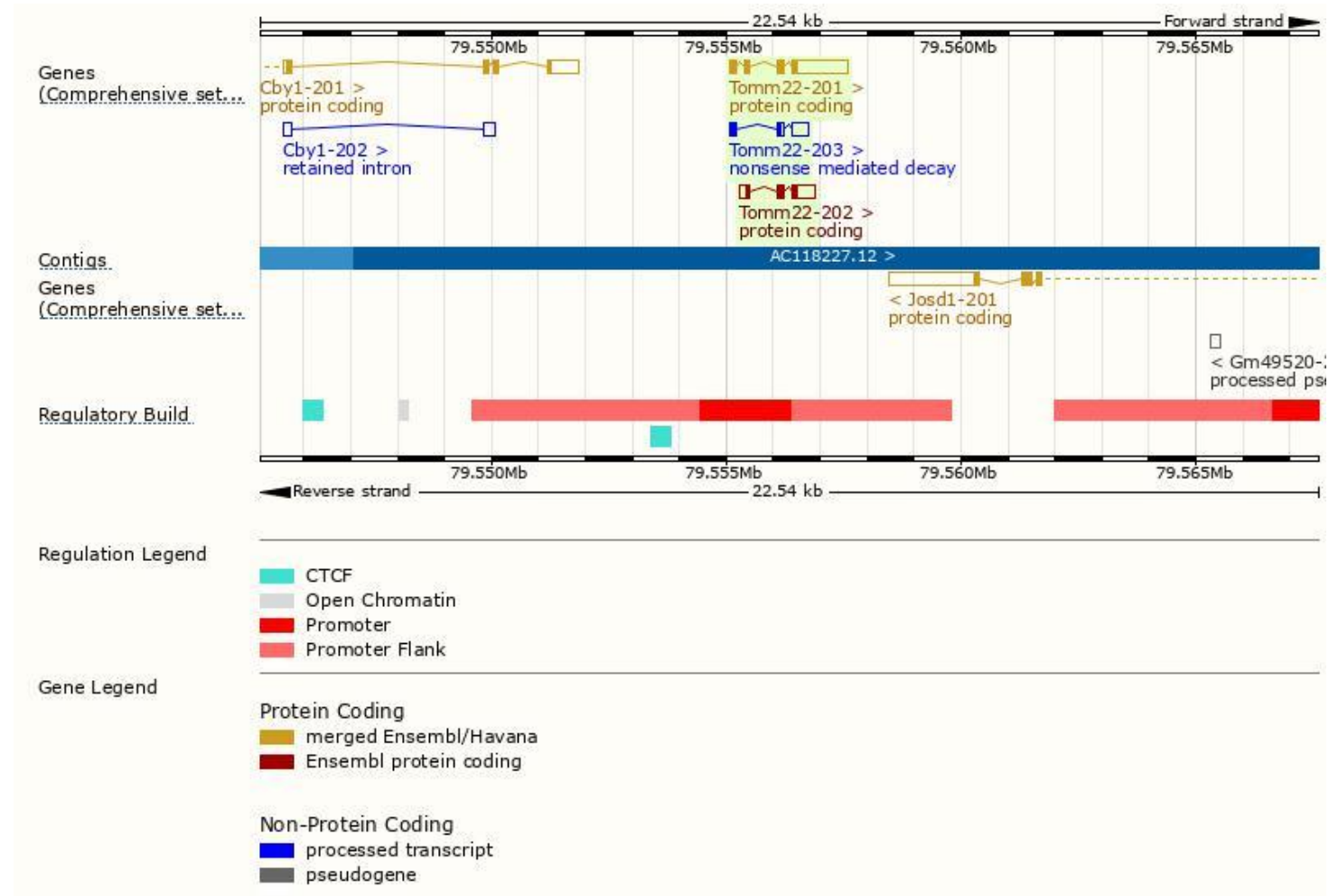
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tomm22-201	<a href="#">ENSMUST00000023062.5</a>	1569	<a href="#">142aa</a>	Protein coding	<a href="#">CCDS27646</a>		TSL:1 , GENCODE basic , APPRIS P2 ,
Tomm22-202	<a href="#">ENSMUST00000127292.2</a>	783	<a href="#">89aa</a>	Protein coding	-		TSL:2 , GENCODE basic , APPRIS ALT2 ,
Tomm22-203	<a href="#">ENSMUST00000229502.2</a>	595	<a href="#">61aa</a>	Nonsense mediated decay	-		

The strategy is based on the design of *Tomm22-201* transcript, the transcription is shown below:



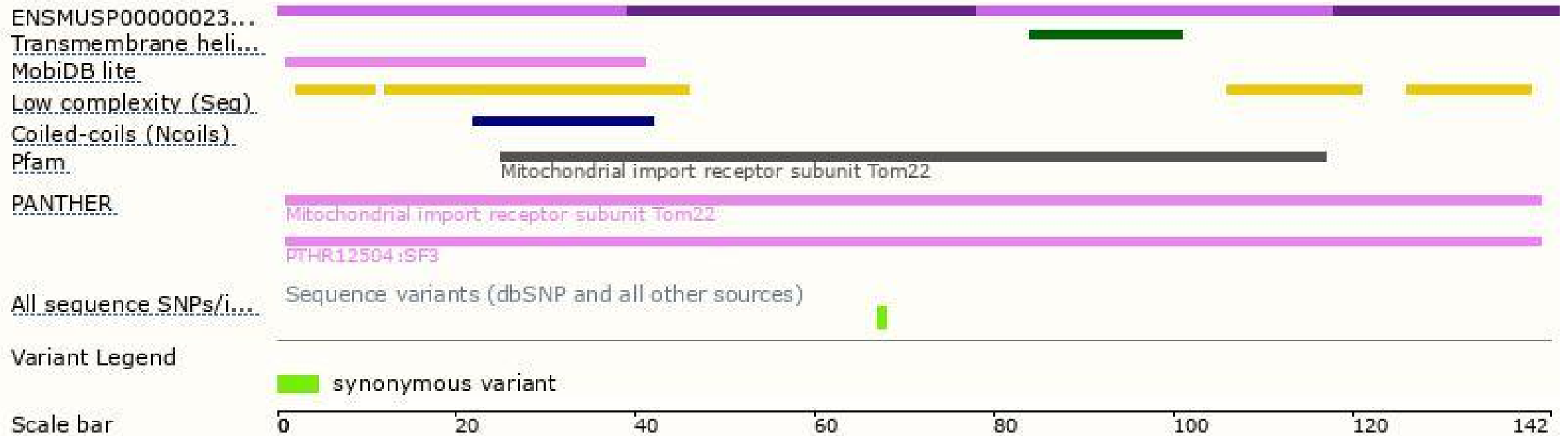
Source: <https://www.ensembl.org>

# Genomic Information



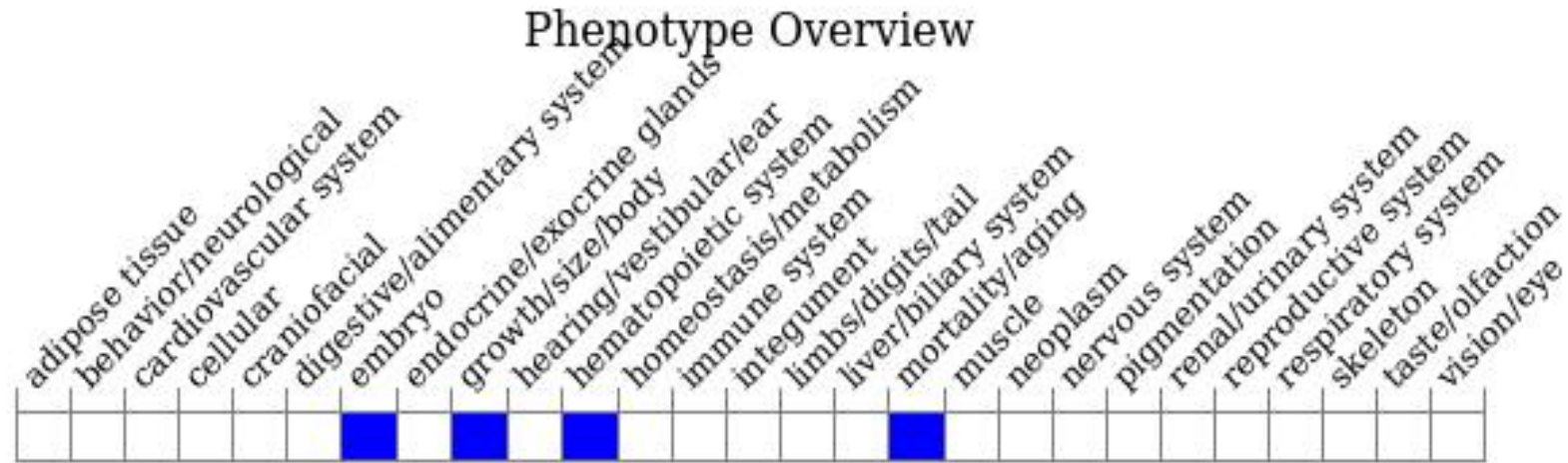


# Protein Information





# Mouse Phenotype Information (MGI)



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

# Important Information

- The effect of *Cby1* and *Josd1* gene is unknown.
- *Tomm22* is located on Chr15. 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.