

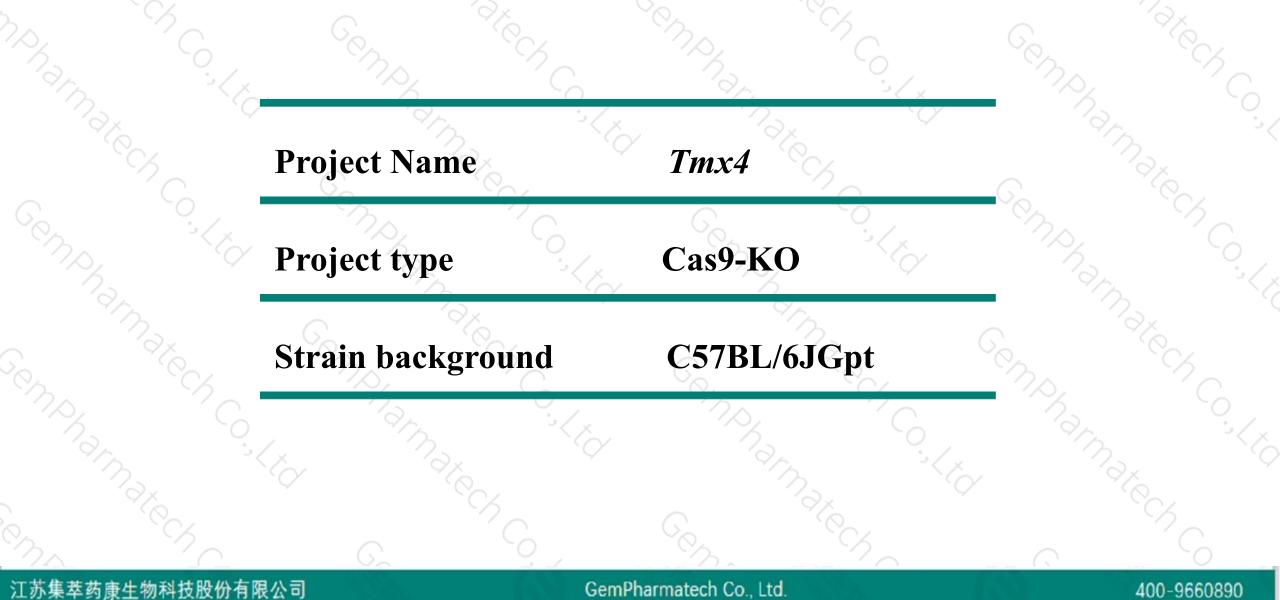
# Tmx4 Cas9-KO Strategy

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### **Project Overview**

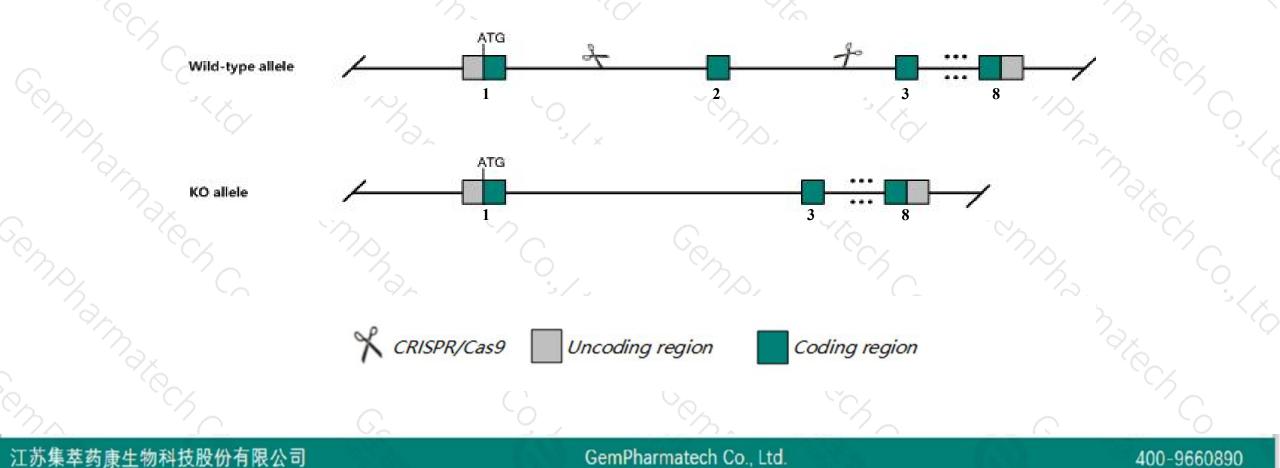




# **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the *Tmx4* gene. The schematic diagram is as follows:





- The *Tmx4* gene has 4 transcripts. According to the structure of *Tmx4* gene, exon2 of *Tmx4-201* (ENSMUST00000038228.10) transcript is recommended as the knockout region. The region contains 116bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Tmx4* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Tmx4* gene is located on the Chr2. 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.

Notice

### **Gene information** (NCBI)



#### Tmx4 thioredoxin-related transmembrane protein 4 [ Mus musculus (house mouse) ]

Gene ID: 52837, updated on 10-Oct-2019

#### Summary

Official Symbol Tmx4 provided by MGI

Official Full Name thioredoxin-related transmembrane protein 4 provided by MGI Primary source

MGI:MGI:106558

Ensembl:ENSMUSG0000034723 See related

Gene type protein coding

RefSeg status PROVISIONAL

Organism Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Lineage Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as Txndc13; Al843224; AW046784; mKIAA1162; D2Bwg1356e; 2810417D04Rik; 4930500L08Rik Expression Broad expression in frontal lobe adult (RPKM 23.2), cerebellum adult (RPKM 18.6) and 21 other tissues See more Orthologs human all

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## **Transcript information (Ensembl)**



#### The gene has 4 transcripts, all transcripts are shown below:

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Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tmx4-201	ENSMUST0000038228.10	5488	<u>335aa</u>	Protein coding	CCDS16784	Q0P5W2 Q8C0L0	TSL:1 GENCODE basic APPRIS P2
Tmx4-203	ENSMUST00000110120.1	2602	<u>183aa</u>	Protein coding	÷	A2ARI0	TSL:1 GENCODE basic APPRIS ALT2
Tmx4-202	ENSMUST00000110119.1	596	<u>166aa</u>	Protein coding	÷	A2ARI1	TSL:2 GENCODE basic
Tmx4-204	ENSMUST00000137377.1	3917	No protein	IncRNA	2	<u>-</u> 4	TSL:1
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The strategy is based on the design of *Tmx4-201* transcript, The transcription is shown below

#### < Tmx4-201 protein coding

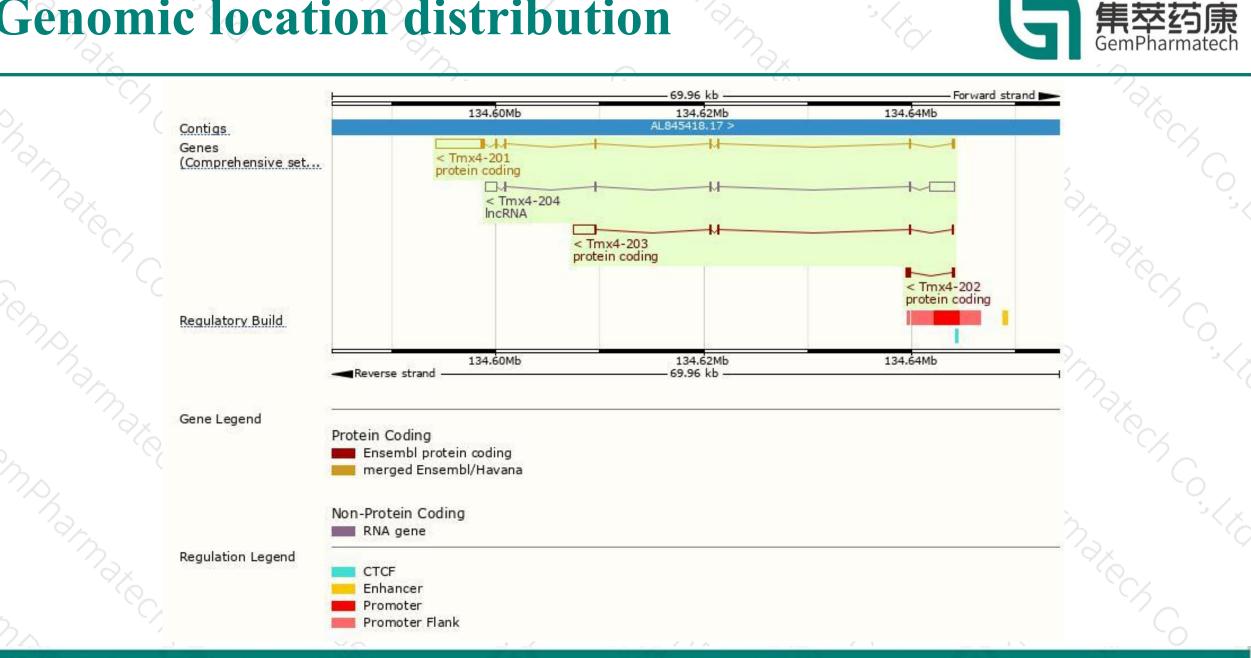
Reverse strand

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49.96 kb

### **Genomic location distribution**

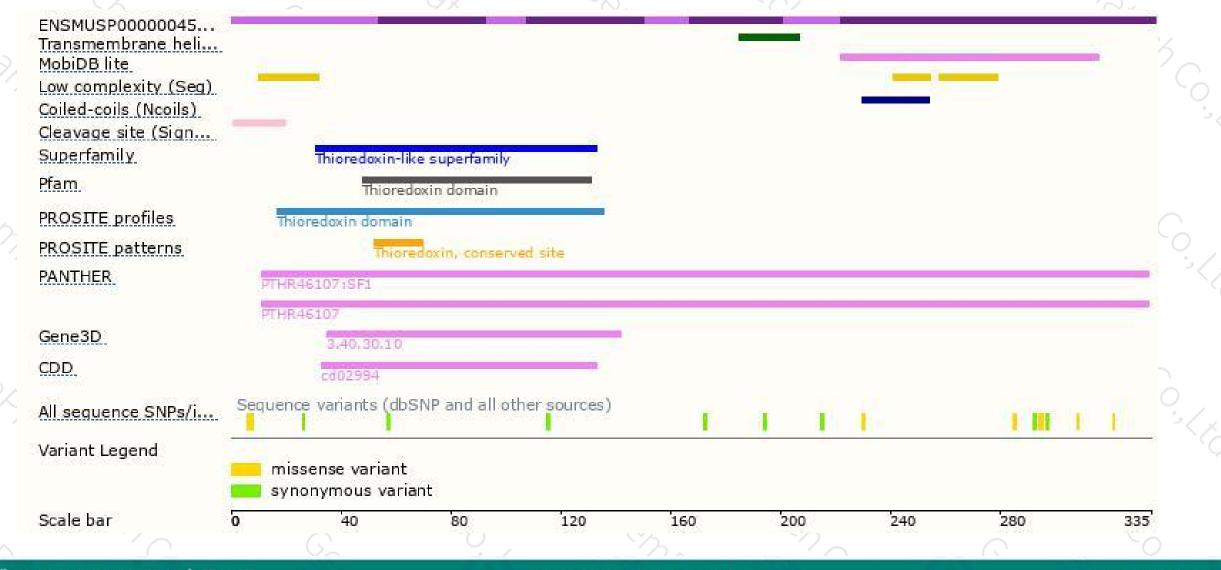


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### **Protein domain**





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If you have any questions, you are welcome to inquire. Tel: 400-9660890



