

Mov10 Cas9-KO Strategy

Designer: Huan Fan

Reviewer: Huan Wang

Design Date: 2019-12-23

Project Overview



Project Name

Mov10

Project type

Cas9-KO

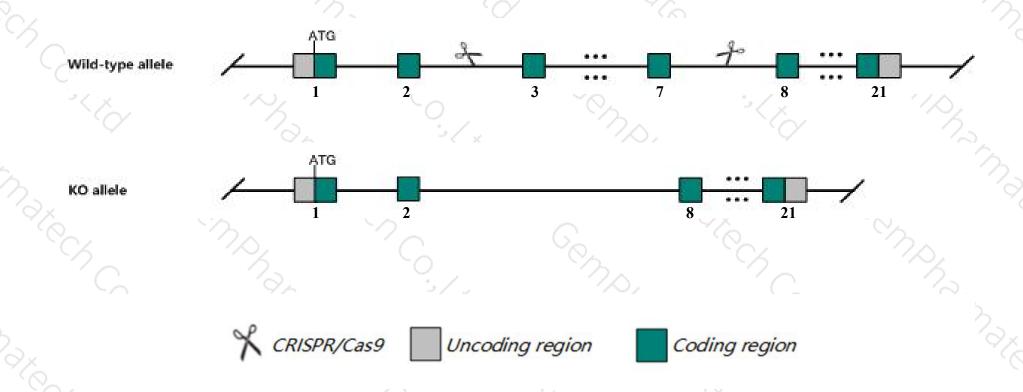
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Mov10* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Mov10* gene has 9 transcripts. According to the structure of *Mov10* gene, exon3-exon7 of *Mov10-206*(ENSMUST00000166979.7) transcript is recommended as the knockout region. The region contains 1006bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Mov10* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ According to the existing MGI data, Homozygous knockout is embryonic lethal. Heterozygous knockout leads to reduced dendritic branching of neurons, which affects anxiety- and/or activity-related behavior.
- > The *Mov10* gene is located on the Chr3. 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)



Mov10 Mov10 RISC complex RNA helicase [Mus musculus (house mouse)]

Gene ID: 17454, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Mov10 provided by MGI

Official Full Name Mov10 RISC complex RNA helicase provided by MGI

Primary source MGI:MGI:97054

See related Ensembl: ENSMUSG00000002227

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 C77703, Mov-10

Expression Ubiquitous expression in duodenum adult (RPKM 17.0), small intestine adult (RPKM 14.9) and 27 other tissuesSee more

Orthologs <u>human</u> all

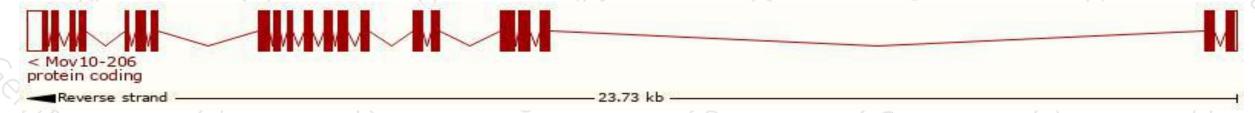
Transcript information (Ensembl)



The gene has 9 transcripts, all transcripts are shown below:

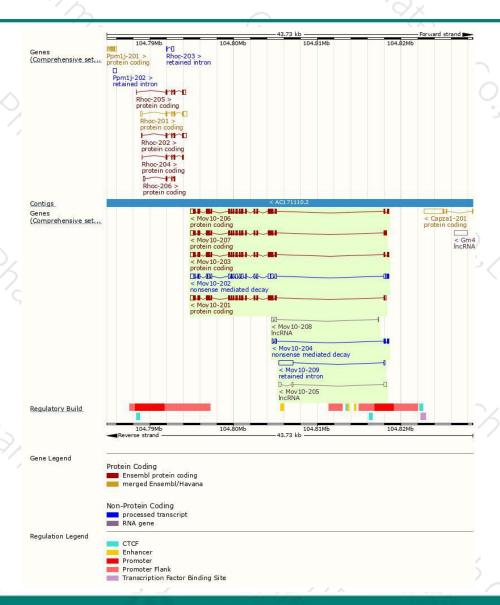
| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|------------|-----------------------|-------|---------------|-------------------------|-----------|---------|---------------------------------|
| Mov10-206 | ENSMUST00000166979.7 | 3584 | <u>1077aa</u> | Protein coding | CCDS51033 | E9PW39 | TSL:1 GENCODE basic |
| Mov10-207 | ENSMUST00000168015.7 | 3460 | 1004aa | Protein coding | CCDS38579 | P23249 | TSL:1 GENCODE basic APPRIS P2 |
| Mov10-203 | ENSMUST00000106775.7 | 3584 | <u>1077aa</u> | Protein coding | ¥ | D3YVL0 | TSL:5 GENCODE basic |
| Mov10-201 | ENSMUST00000002297.11 | 3498 | 1004aa | Protein coding | 2 | D3Z3E8 | TSL:5 GENCODE basic APPRIS ALT1 |
| Mov10-202 | ENSMUST00000106774.7 | 3718 | <u>57aa</u> | Nonsense mediated decay | | D3YVL1 | TSL:5 |
| Mov10-204 | ENSMUST00000136148.2 | 755 | <u>145aa</u> | Nonsense mediated decay | - | D3YYR1 | TSL:3 |
| /lov10-209 | ENSMUST00000196211.1 | 1876 | No protein | Retained intron | ¥ | - | TSL:1 |
| Mov10-205 | ENSMUST00000145414.1 | 763 | No protein | IncRNA | 2 | 1028 | TSL:2 |
| Mov10-208 | ENSMUST00000196161.1 | 454 | No protein | IncRNA | | 1871 | TSL:2 |
| | | 7 7 7 | | | W 1000 | 1 | |

The strategy is based on the design of Mov10-206 transcript, The transcription is shown below



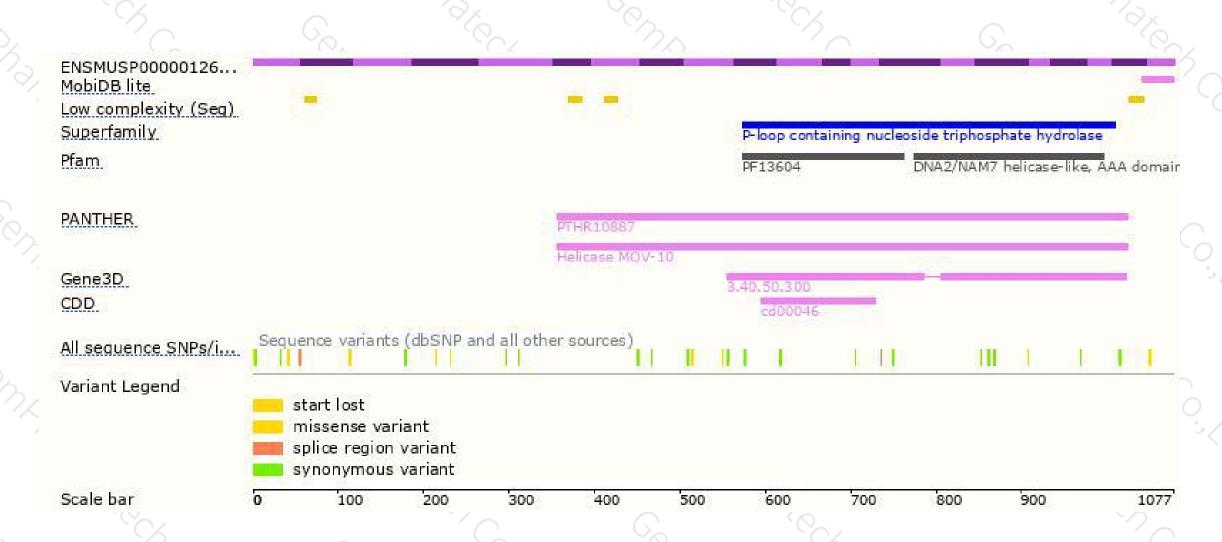
Genomic location distribution





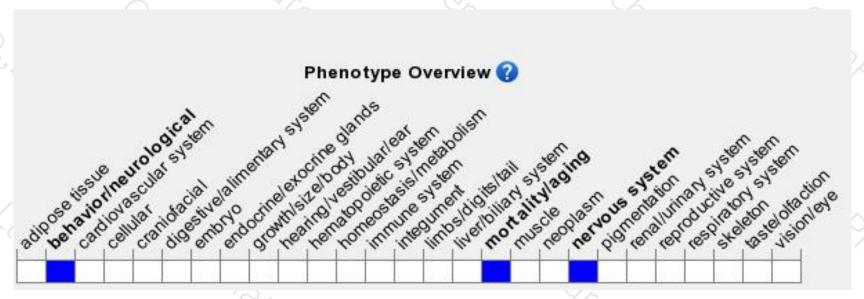
Protein domain





Mouse phenotype description(MGI)





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

According to the existing MGI data, Homozygous knockout is embryonic lethal. Heterozygous knockout leads to reduced dendritic branching of neurons, which affects anxiety- and/or activity-related behavior.



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





