

Musk Cas9-KO Strategy

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Project Overview



Project Name

Musk

Project type

Cas9-KO

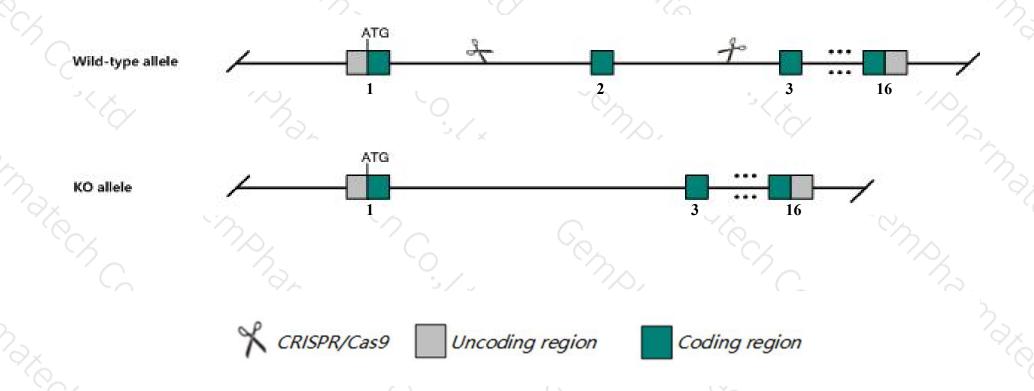
Strain background

C57BL/6JGpt

Knockout strategy



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



Technical routes



The Musk gene has 8 transcripts. According to the structure of Musk gene, exon2 of Musk-203

(ENSMUST00000098057.9) transcript is recommended as the knockout region. The region contains 127bp coding sequence.

Knock out the region will result in disruption of protein function.

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

Notice



According to the existing MGI data, Homozygotes for a targeted null mutation lack neuromuscular synapses and spontaneous movement, and die at birth of respiratory failure.

The *Musk* gene is located on the Chr4. 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



Musk muscle, skeletal, receptor tyrosine kinase [Mus musculus (house mouse)]

Gene ID: 18198, updated on 26-Mar-2019

Summary

☆ ?

Official Symbol Musk provided by MGI

Official Full Name muscle, skeletal, receptor tyrosine kinase provided by MGI

Primary source MGI:MGI:103581

See related Ensembl: ENSMUSG00000057280

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as Mdk4, Mlk, Nsk1, Nsk2, Nsk3

Summary This gene encodes a member of the protein tyrosine kinase family. The encoded protein is a type 1 receptor-like protein located in muscle

membrane that is activated by the heparan sulfate proteoglycan agrin released by nerve cells. The encoded protein activates signaling cascades responsible for multiple aspects of motor neuron and muscle development, including organization of the postsynaptic membrane, synaptic gene transcription, patterning of skeletal muscle, anchoring of acetylcholinesterase, and guidance of motor axons. Alternatively

spliced transcript variants encoding different isoforms have been described. [provided by RefSeq, Jul 2008]

Expression Broad expression in limb E14.5 (RPKM 1.2), colon adult (RPKM 0.4) and 16 other tissuesSee more

Orthologs <u>human</u> <u>all</u>

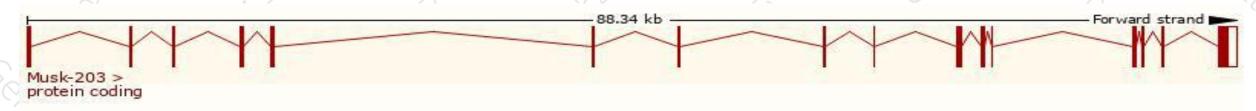
Transcript information Ensembl



The gene has 8 transcripts, all transcripts are shown below:

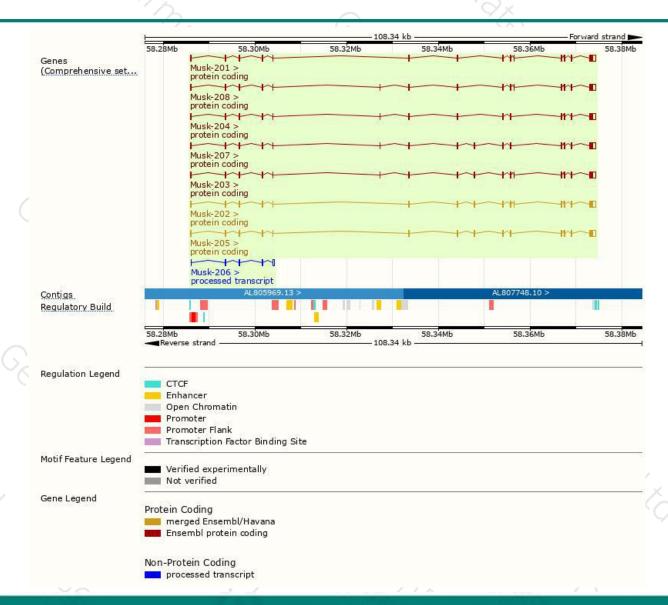
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Musk-203	ENSMUST00000098057.9	3422	893aa	Protein coding	CCDS51182	E9PVV8	TSL:1 GENCODE basic
Musk-207	ENSMUST00000177951.7	3352	<u>870aa</u>	Protein coding	CCDS38770	Q497X0	TSL:5 GENCODE basic APPRIS ALT2
Musk-202	ENSMUST00000084578.11	3347	868aa	Protein coding	CCDS18211	Q32S50 Q61006	TSL:5 GENCODE basic APPRIS P4
Musk-205	ENSMUST00000102893.9	3323	860aa	Protein coding	CCDS18210	Q32S49 Q61006	TSL:5 GENCODE basic APPRIS ALT2
Musk-208	ENSMUST00000179951.1	3376	<u>878aa</u>	Protein coding		J3QPS3	TSL:5 GENCODE basic APPRIS ALT2
Musk-204	ENSMUST00000098059.9	3352	<u>870aa</u>	Protein coding		F6WVU1	TSL:5 GENCODE basic APPRIS ALT2
Musk-201	ENSMUST00000081919.11	3348	<u>868aa</u>	Protein coding	11	K3W4P7	TSL:1 GENCODE basic APPRIS ALT2
Musk-206	ENSMUST00000135879.1	903	No protein	Processed transcript	12	22	TSL:3

The strategy is based on the design of *Musk-203* transcript, The transcription is shown below



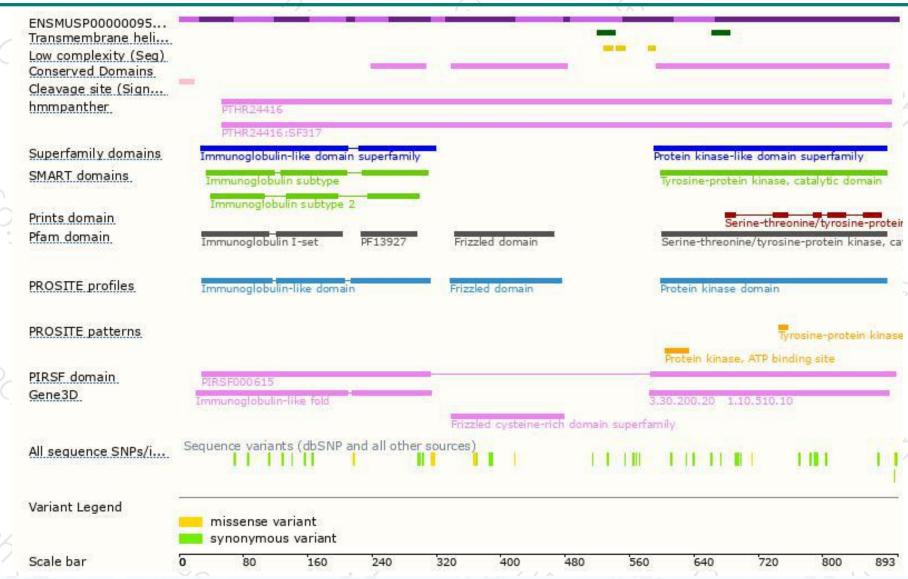
Genomic location distribution





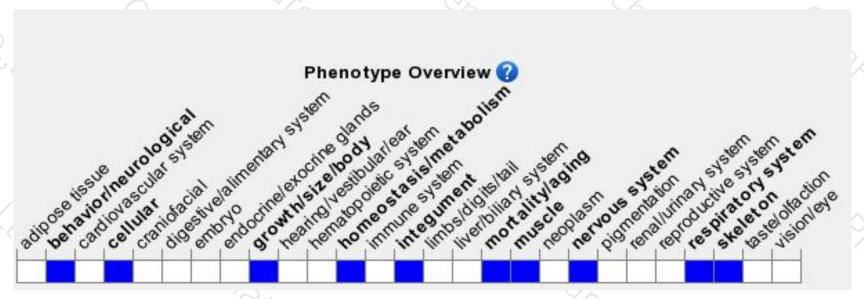
Protein domain





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, Homozygotes for a targeted null mutation lack neuromuscular synapses and spontaneous movement, and die at birth of respiratory failure.



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





