

# *Myh3* Cas9-CKO Strategy

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# Project Overview

**Project Name**

*Myh3*

**Project type**

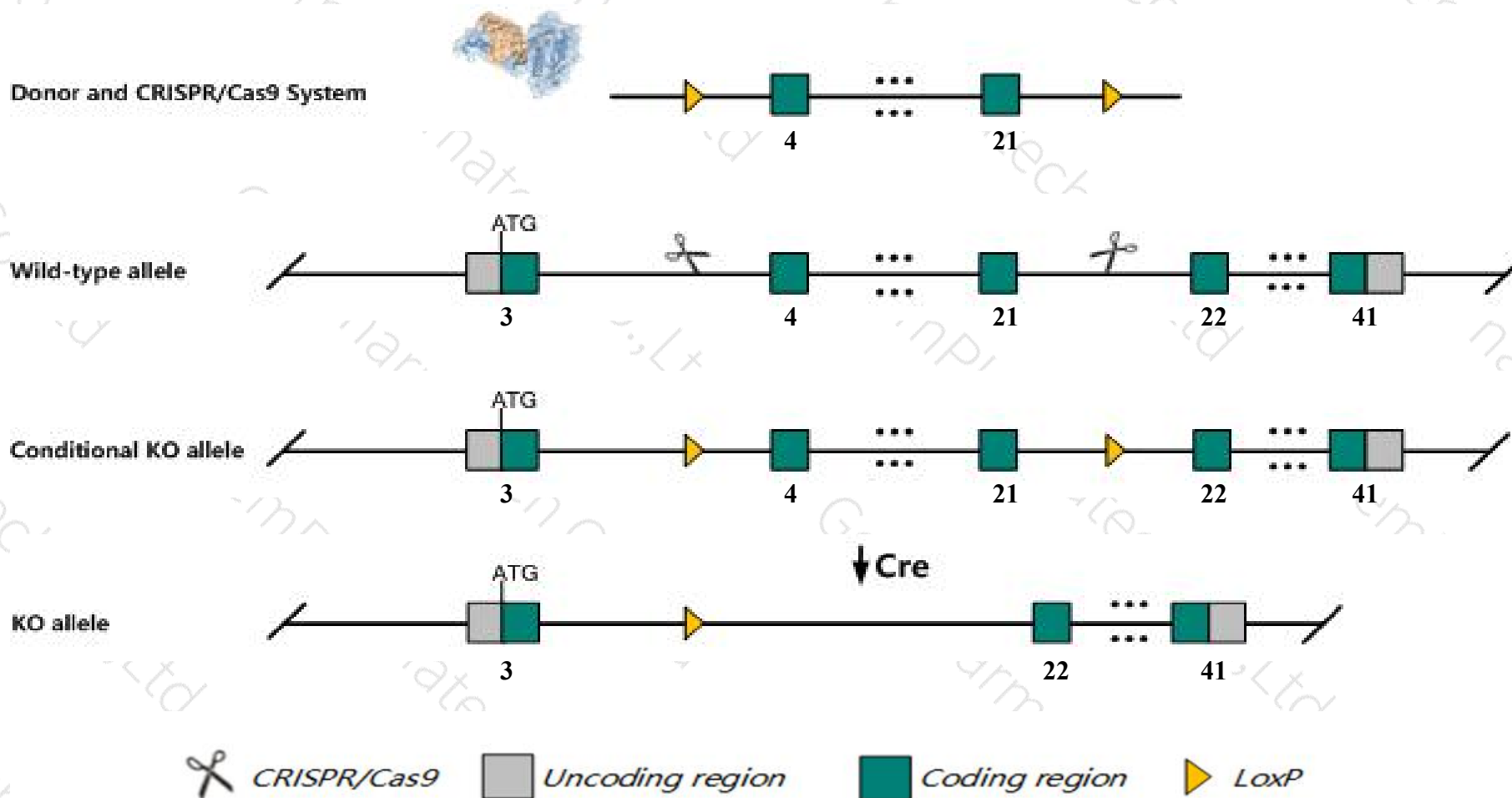
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Myh3* gene. The schematic diagram is as follows:



- The *Myh3* gene has 3 transcripts. According to the structure of *Myh3* gene, exon4-exon21 of *Myh3-202* (ENSMUST00000108689.7) transcript is recommended as the knockout region. The region contains 2222bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Myh3* gene. The brief process is as follows: CRISPR/Cas9 system and Donor 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 sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

# Notice

- The floxed region is near to the N-terminal of *Mir6923* gene, this strategy may influence the regulatory function of the N-terminal of *Mir6923* gene.
- The *Myh3* gene is located on the Chr11. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

# Gene information (NCBI)

## Myh3 myosin, heavy polypeptide 3, skeletal muscle, embryonic [ *Mus musculus* (house mouse) ]

Gene ID: 17883, updated on 21-Aug-2019

### Summary

<b>Official Symbol</b>	Myh3 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	myosin, heavy polypeptide 3, skeletal muscle, embryonic provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1339709</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000020908</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	REVIEWED
<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>	Myhse; Myhs-e; MyHC-emb
<b>Summary</b>	Myosin is a major contractile protein which converts chemical energy into mechanical energy through the hydrolysis of ATP. Myosin is a hexameric protein composed of a pair of myosin heavy chains (MYH) and two pairs of nonidentical light chains. This gene is a member of the MYH family and encodes a protein with an IQ domain and a myosin head-like domain. [provided by RefSeq, Sep 2015]
<b>Expression</b>	Biased expression in limb E14.5 (RPKM 52.1), CNS E14 (RPKM 5.2) and 1 other tissue <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

### Genomic context

**Location:** 11 B3; 11 40.59 cM

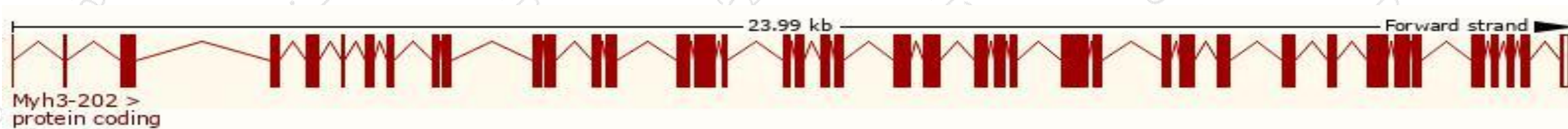
See Myh3 in [Genome Data Viewer](#)

# Transcript information (Ensembl)

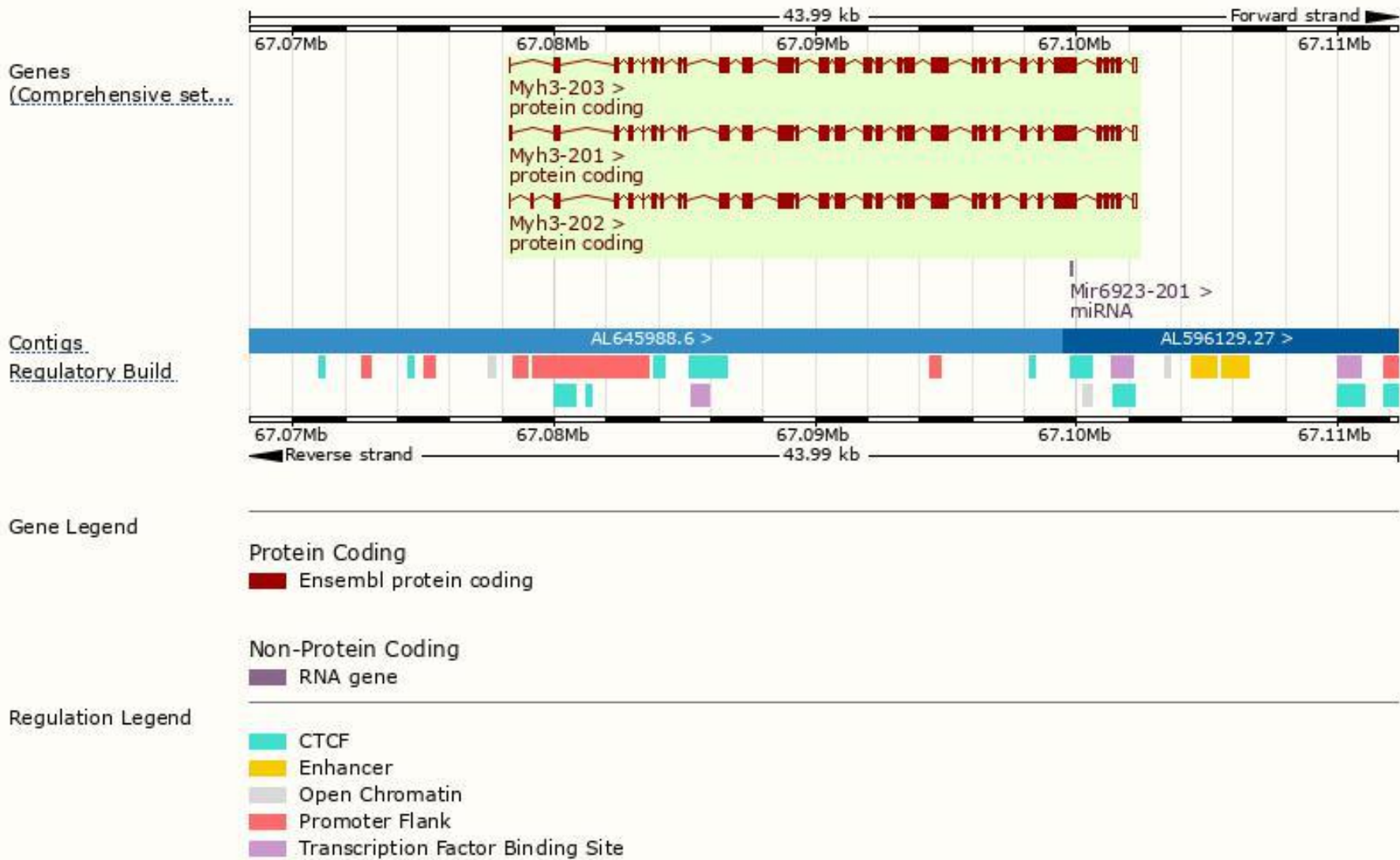
The gene has 3 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Myh3-202	<a href="#">ENSMUST00000108689.7</a>	6031	<a href="#">1940aa</a>	Protein coding	<a href="#">CCDS36184</a>	<a href="#">P13541</a>	TSL:5 GENCODE basic APPRIS P1
Myh3-203	<a href="#">ENSMUST00000165221.1</a>	5994	<a href="#">1940aa</a>	Protein coding	<a href="#">CCDS36184</a>	<a href="#">P13541</a>	TSL:5 GENCODE basic APPRIS P1
Myh3-201	<a href="#">ENSMUST00000007301.13</a>	5992	<a href="#">1940aa</a>	Protein coding	<a href="#">CCDS36184</a>	<a href="#">P13541</a>	TSL:5 GENCODE basic APPRIS P1

The strategy is based on the design of *Myh3-202* transcript, The transcription is shown below



# Genomic location distribution





# Protein domain



If you have any questions, you are welcome to inquire.

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