

Myoz1 Cas9-CKO Strategy

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Project Overview

Project Name

Myoz1

Project type

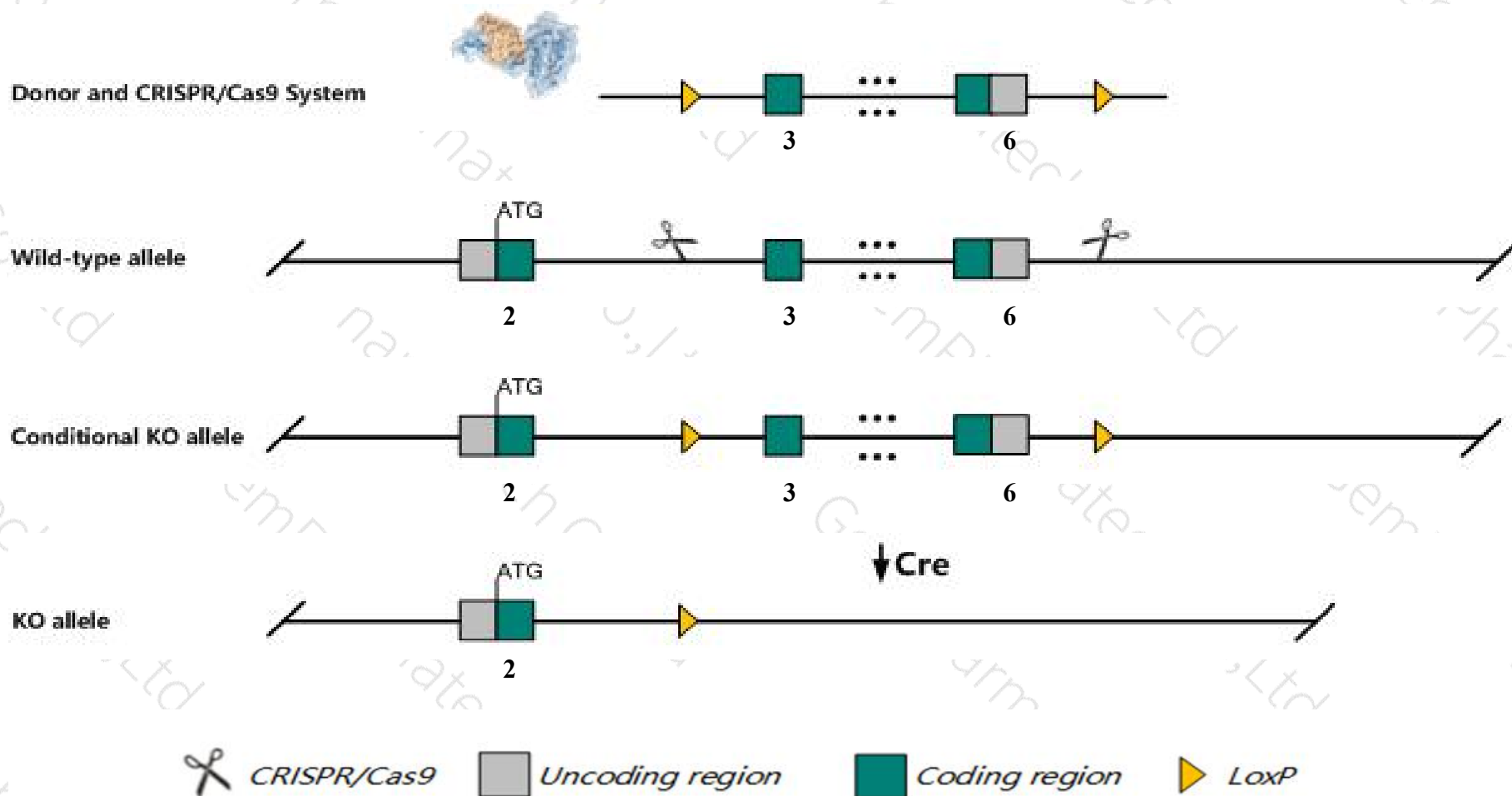
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Myoz1* gene. The schematic diagram is as follows:



Technical routes

- The *Myoz1* gene has 4 transcripts. According to the structure of *Myoz1* gene, exon3-exon6 of *Myoz1-201* (ENSMUST00000090469.7) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Myoz1* 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.

- According to the existing MGI data, Mice homozygous for a knock-out allele show reduced body weight and fast-twitch muscle mass, a fiber type shift toward more oxidative fibers, increased exercise capacity and calcineurin activity, and enhanced muscle regeneration after cardiotoxin injury.
- The *Myoz1* gene is located on the Chr14. 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)

Myoz1 myozenin 1 [*Mus musculus* (house mouse)]

Gene ID: 59011, updated on 12-Aug-2019

Summary

- Official Symbol

Myoz1 provided by [MGI](#)
- Official Full Name

myozenin 1 provided by [MGI](#)
- Primary source

[MGI:MGI:1929471](#)
- See related

[Ensembl:ENSMUSG00000068697](#)
- 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

FATZ; Myoz; AV090278; 2310001N11Rik
- Expression

Biased expression in mammary gland adult (RPKM 65.7) and lung adult (RPKM 3.9) [See more](#)
- Orthologs

[human](#) [all](#)

Genomic context

Location: 14; 14 A3

See Myoz1 in [Genome Data Viewer](#)

Exon count: 6

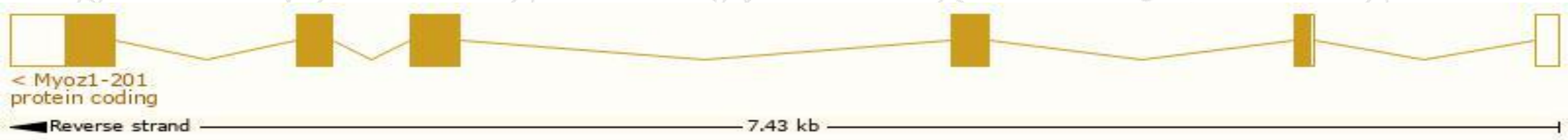
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	14	NC_000080.6 (20649102..20656540, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	14	NC_000080.5 (21468324..21475762, complement)

Transcript information (Ensembl)

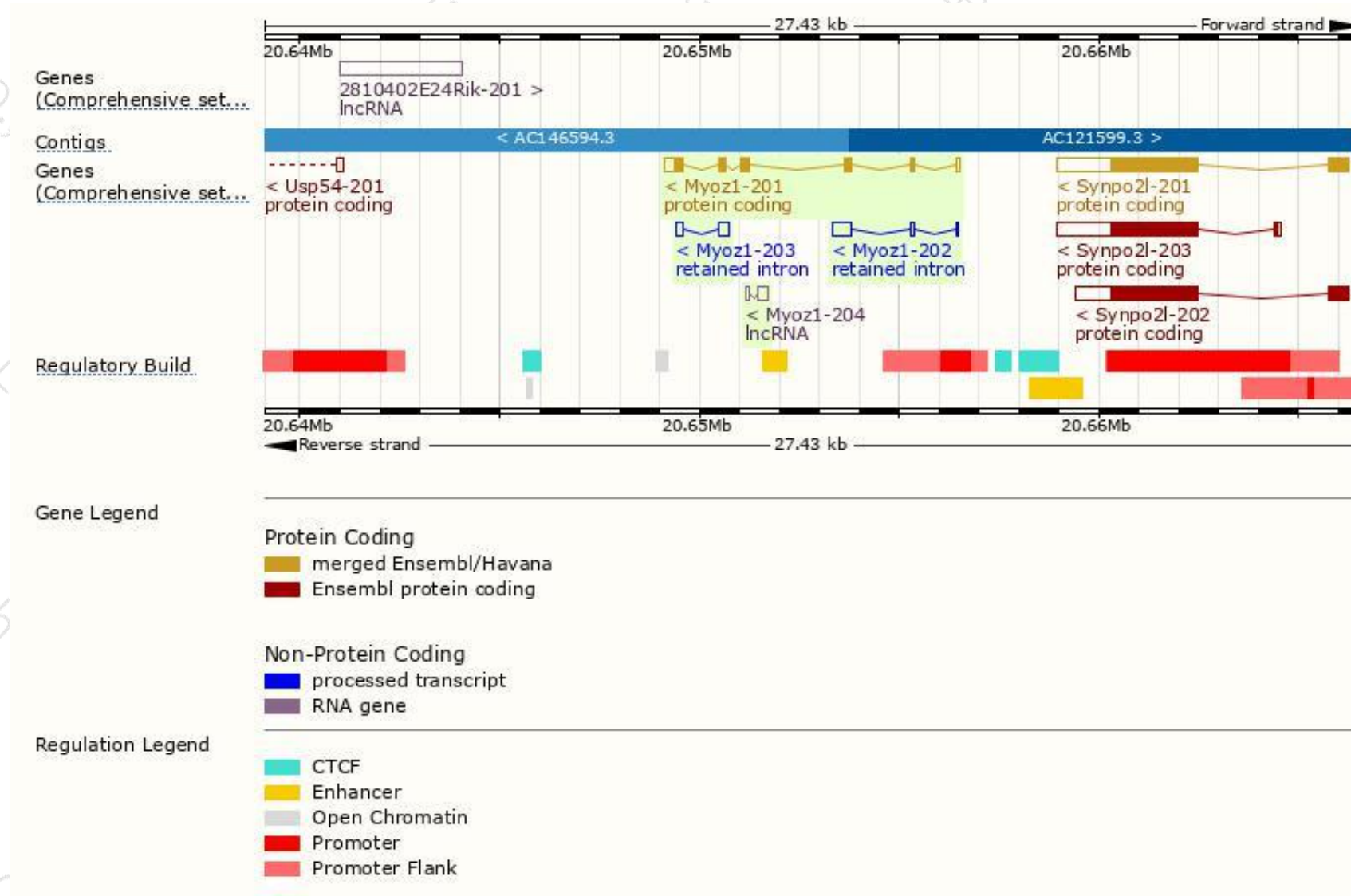
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Myoz1-201	ENSMUST00000090469.7	1289	296aa	Protein coding	CCDS36819	Q9JK37	TSL:1 GENCODE basic APPRIS P1
Myoz1-202	ENSMUST00000224436.1	607	No protein	Retained intron	-	-	
Myoz1-203	ENSMUST00000224472.1	405	No protein	Retained intron	-	-	
Myoz1-204	ENSMUST00000225231.1	346	No protein	lncRNA	-	-	

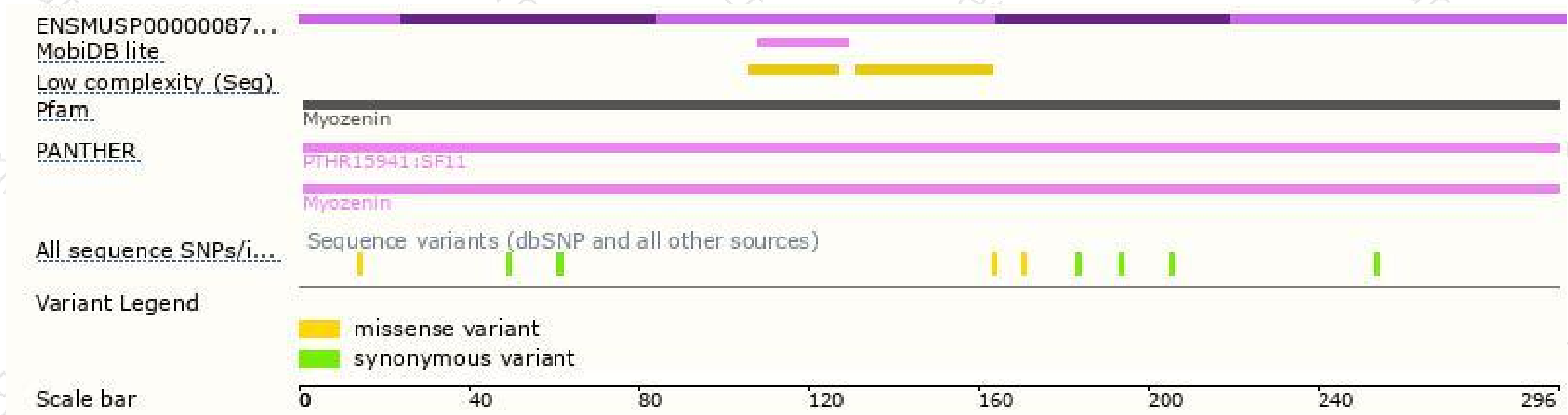
The strategy is based on the design of *Myoz1-201* transcript,The transcription is shown below



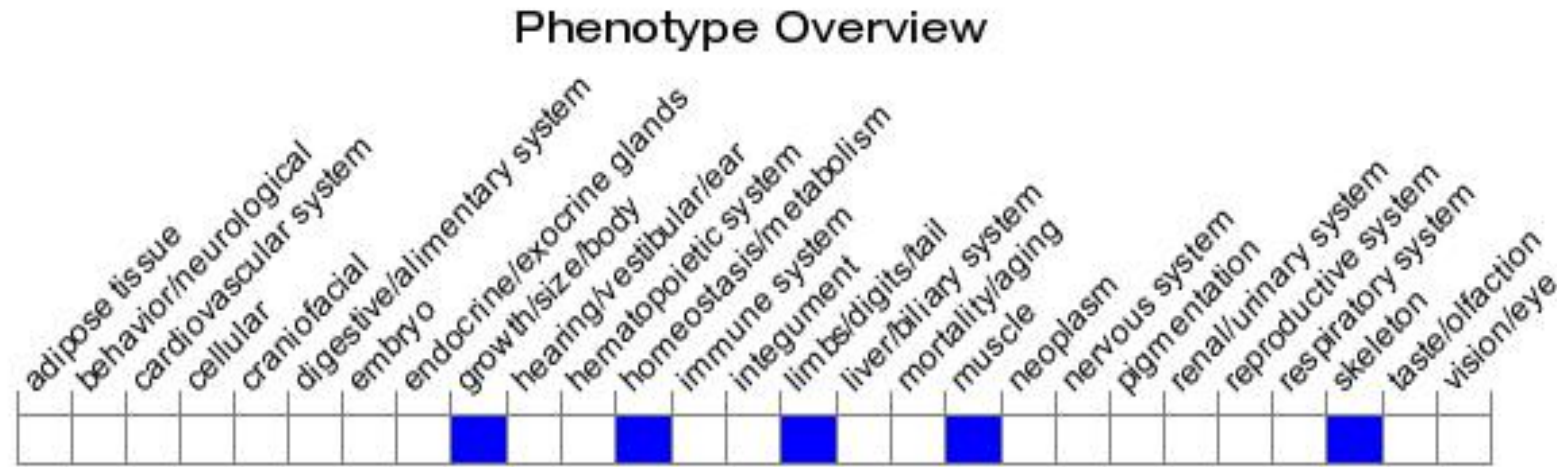
Genomic location distribution



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, Mice homozygous for a knock-out allele show reduced body weight and fast-twitch muscle mass, a fiber type shift toward more oxidative fibers, increased exercise capacity and calcineurin activity, and enhanced muscle regeneration after cardiotoxin injury.

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

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