

# Myoz3 Cas9-KO Strategy

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# **Project Overview**



Project Name Myoz3

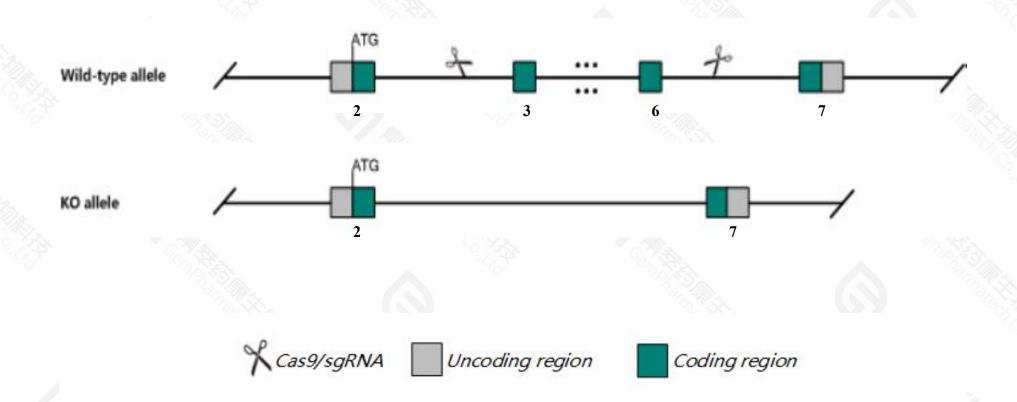
Project type Cas9-KO

Strain background C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Myoz3 gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The *Myoz3* gene has 1 transcript. According to the structure of *Myoz3* gene, exon3-exon6 of *Myoz3-201*(ENSMUST00000056533.9) transcript is recommended as the knockout region. The region contains 508bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Myoz3* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA 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.

### **Notice**



- > The *Myoz3* gene is located on the Chr18. 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)



#### Myoz3 myozenin 3 [Mus musculus (house mouse)]

Gene ID: 170947, updated on 25-Sep-2020

#### Summary

☆ ?

Official Symbol Myoz3 provided by MGI

Official Full Name myozenin 3 provided by MGI

Primary source MGI:MGI:2179296

See related Ensembl:ENSMUSG00000049173

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 4833419K08Rik, Fatz-, Fatz-3, cals

Expression Biased expression in mammary gland adult (RPKM 4.7), lung adult (RPKM 0.8) and 1 other tissueSee more

Orthologs <u>human all</u>

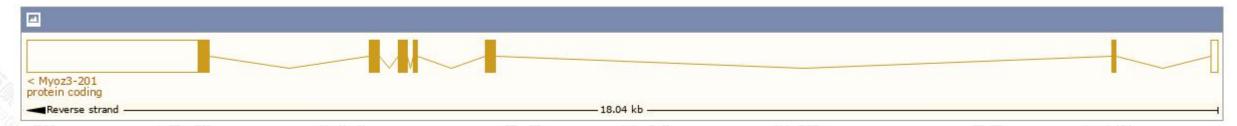
# Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

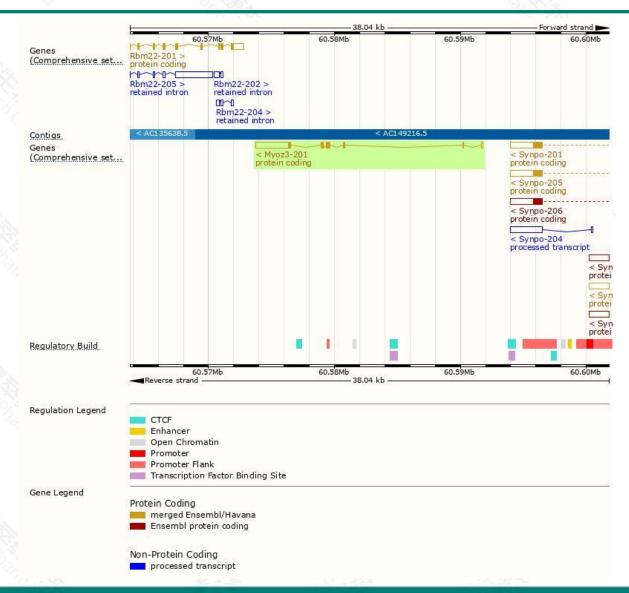
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Myoz3-201	ENSMUST00000056533.9	3436	245aa	Protein coding	CCDS29273		TSL:1 , GENCODE basic , APPRIS P1 ,

The strategy is based on the design of *Myoz3-201* transcript, the transcription is shown below:



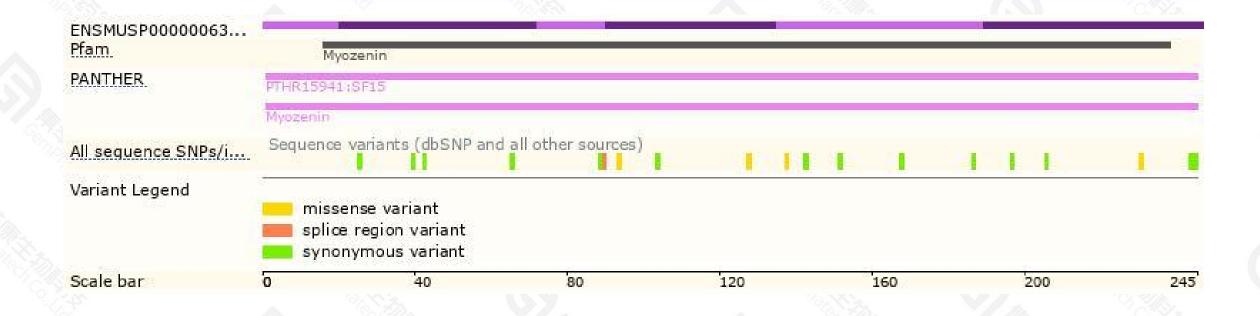
### Genomic location distribution





### Protein domain







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

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