

Myl9 Cas9-CKO Strategy

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Project Overview

Project Name

Myl9

Project type

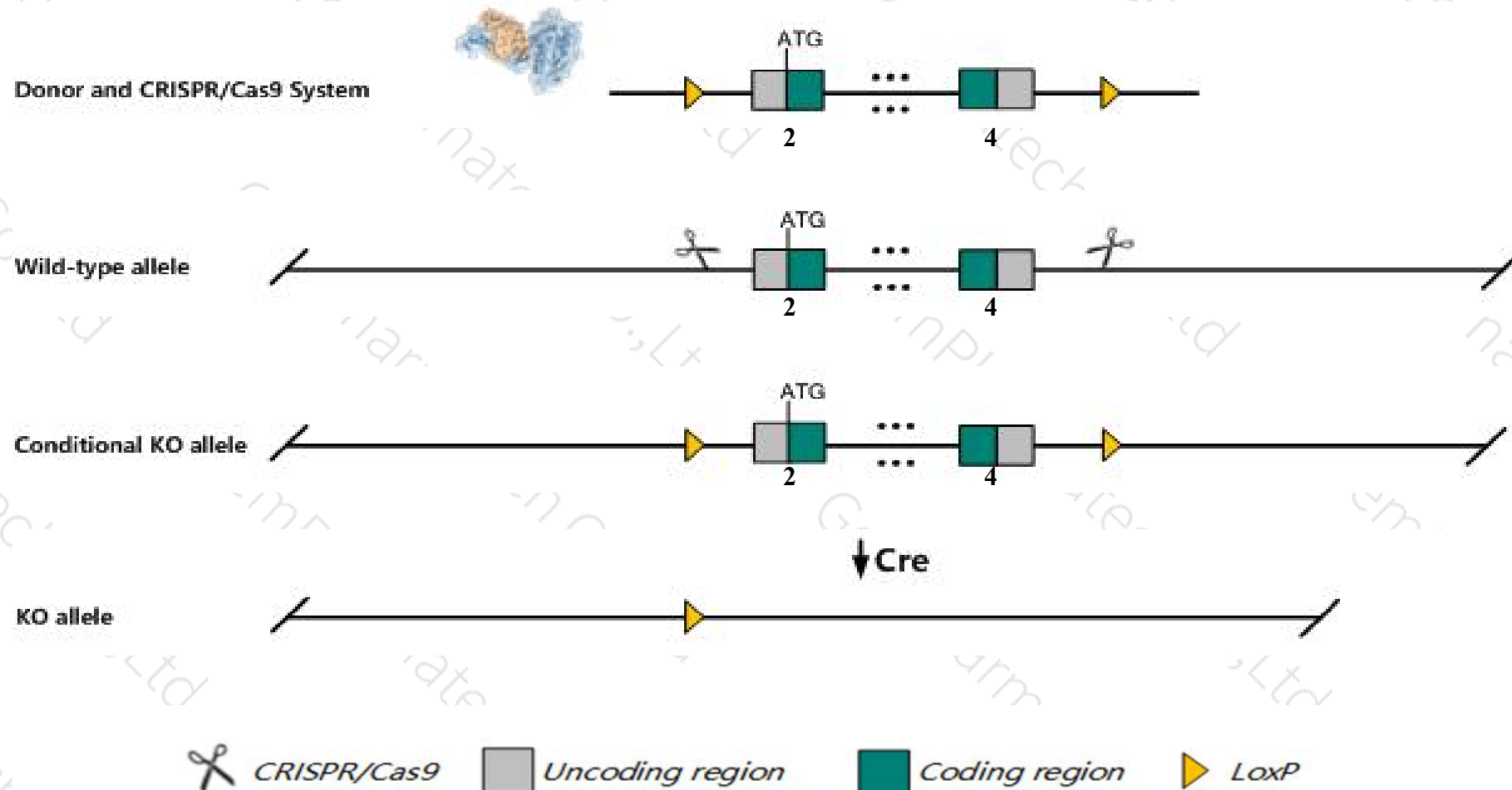
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Myf9* gene. The schematic diagram is as follows:



- The *Myl9* gene has 4 transcripts. According to the structure of *Myl9* gene, exon2-exon4 of *Myl9-201* (ENSMUST00000088552.6) transcript is recommended as the knockout region. The region contains all 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 *Myl9* 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 *Myl9* gene is located on the Chr2. 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)

Myl9 myosin, light polypeptide 9, regulatory [*Mus musculus* (house mouse)]

Gene ID: 98932, updated on 26-Nov-2019

Summary

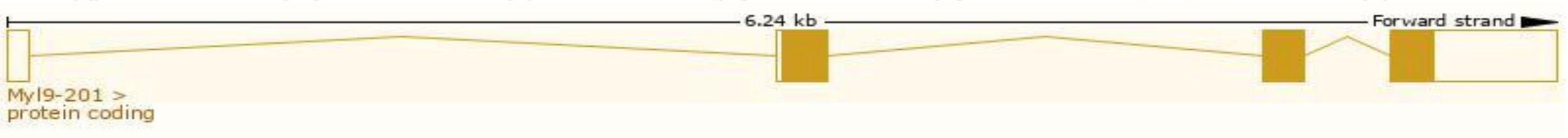
Official Symbol	Myl9 provided by MGI
Official Full Name	myosin, light polypeptide 9, regulatory provided by MGI
Primary source	MGI:MGI:2138915
See related	Ensembl:ENSMUSG000000067818
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	MLC20; RLC-C; Mylc2c; AI327049
Expression	Biased expression in bladder adult (RPKM 1647.7), colon adult (RPKM 313.1) and 8 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

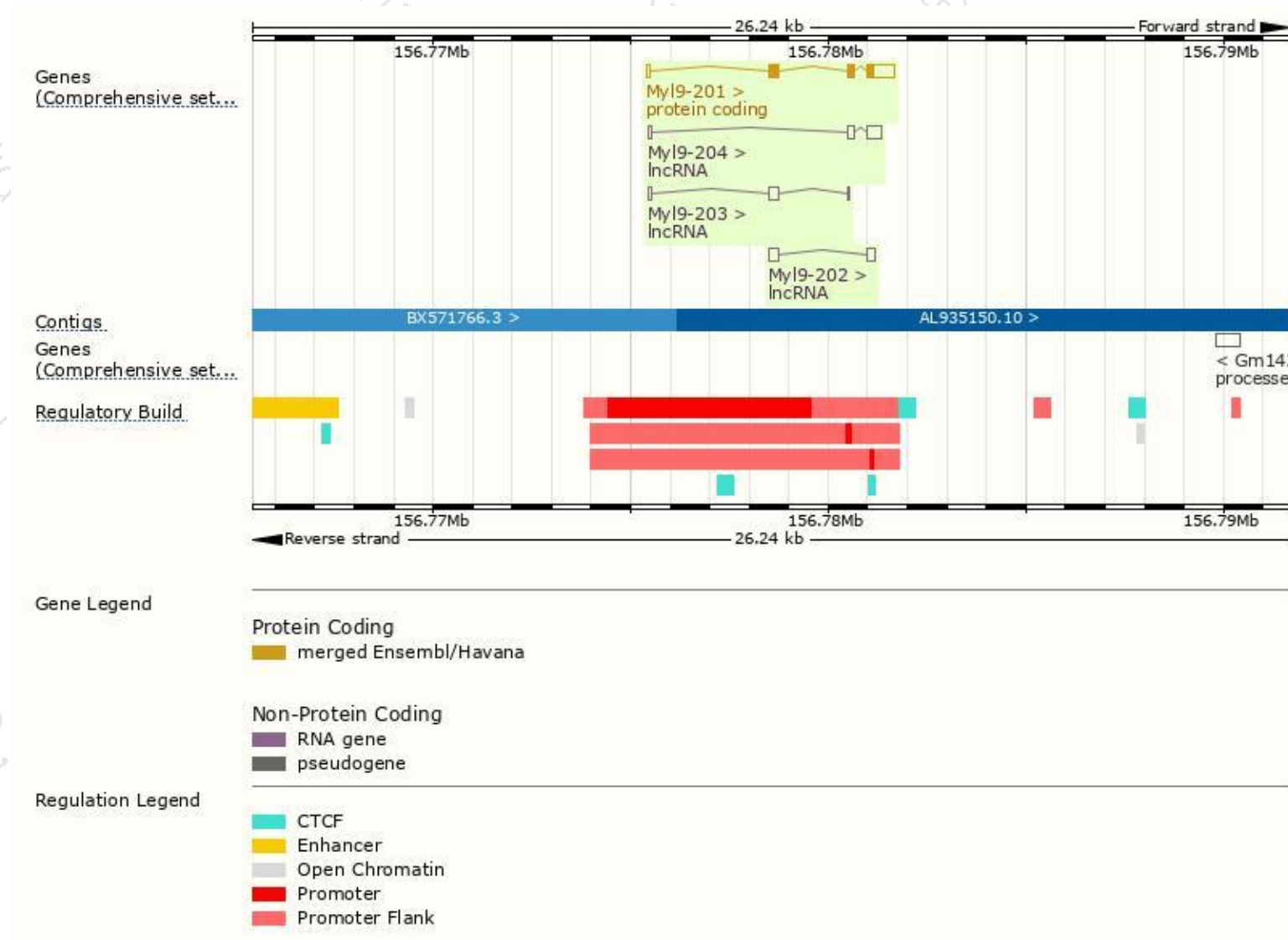
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
MyI9-201	ENSMUST00000088552.6	1128	172aa	Protein coding	CCDS50779	Q9CQ19	TSL:1 GENCODE basic APPRIS P1
MyI9-204	ENSMUST00000145299.1	619	No protein	lncRNA	-	-	TSL:3
MyI9-202	ENSMUST00000131622.1	387	No protein	lncRNA	-	-	TSL:5
MyI9-203	ENSMUST00000132985.1	345	No protein	lncRNA	-	-	TSL:3

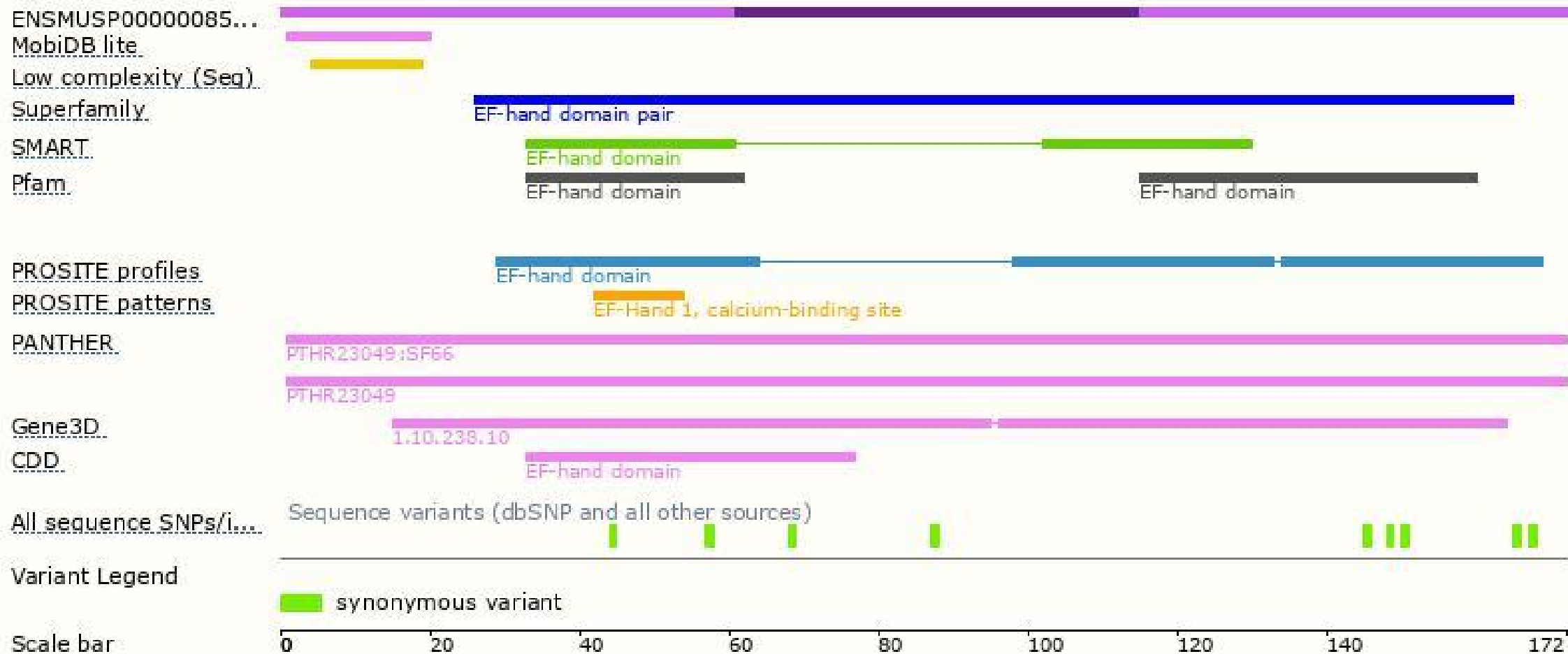
The strategy is based on the design of *MyI9-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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