

Fabp3 Cas9-CKO Strategy

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Project Overview

Project Name

Fabp3

Project type

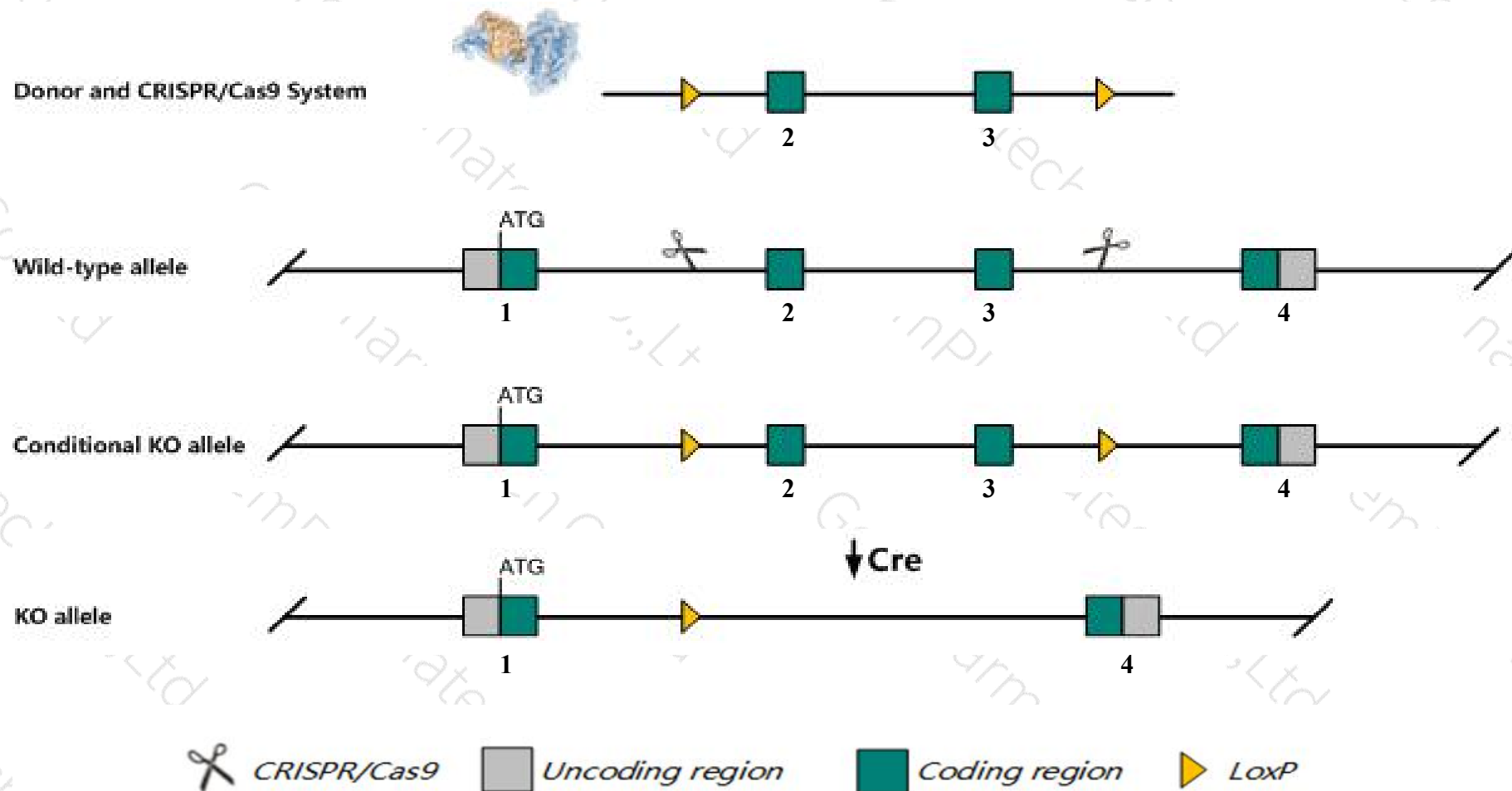
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fabp3* gene. The schematic diagram is as follows:



- The *Fabp3* gene has 1 transcript. According to the structure of *Fabp3* gene, exon2-exon3 of *Fabp3-201* (ENSMUST00000070532.7) transcript is recommended as the knockout region. The region contains 275bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fabp3* 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, Inactivation of this locus results in impaired fatty acid utilization. Homozygous null mice show exercise intolerance and cardiac hypertrophy.
- The floxed region is near to the N-terminal of *Gm10570* and C-terminal of *Zcchc17* gene, this strategy may influence the regulatory function of the N-terminal of *Gm10570* and C-terminal of *Zcchc17* gene.
- The *Fabp3* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Fabp3 fatty acid binding protein 3, muscle and heart [*Mus musculus* (house mouse)]

Gene ID: 14077, updated on 12-Oct-2019

Summary

Official Symbol	Fabp3 provided by MGI
Official Full Name	fatty acid binding protein 3, muscle and heart provided by MGI
Primary source	MGI:MGI:95476
See related	Ensembl:ENSMUSG00000028773
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	Mdgi; Fabph1; Fabph4; H-FABP; Fabph-1; Fabph-4
Expression	Biased expression in heart adult (RPKM 2334.3) and placenta adult (RPKM 178.9) See more
Orthologs	human all

Genomic context

Location: 4 D2.2; 4 63.43 cM

[See Fabp3 in Genome Data Viewer](#)

Exon count: 4

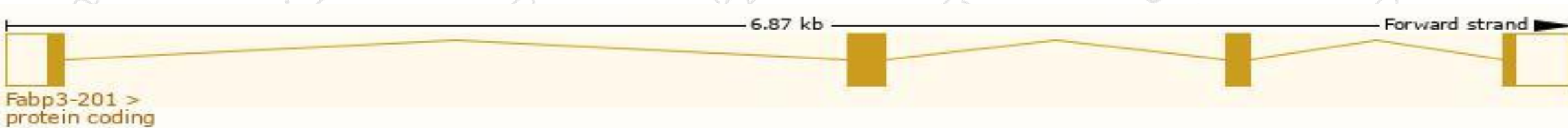
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	4	NC_000070.6 (130308738..130315463)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	4	NC_000070.5 (129986022..129992707)

Transcript information (Ensembl)

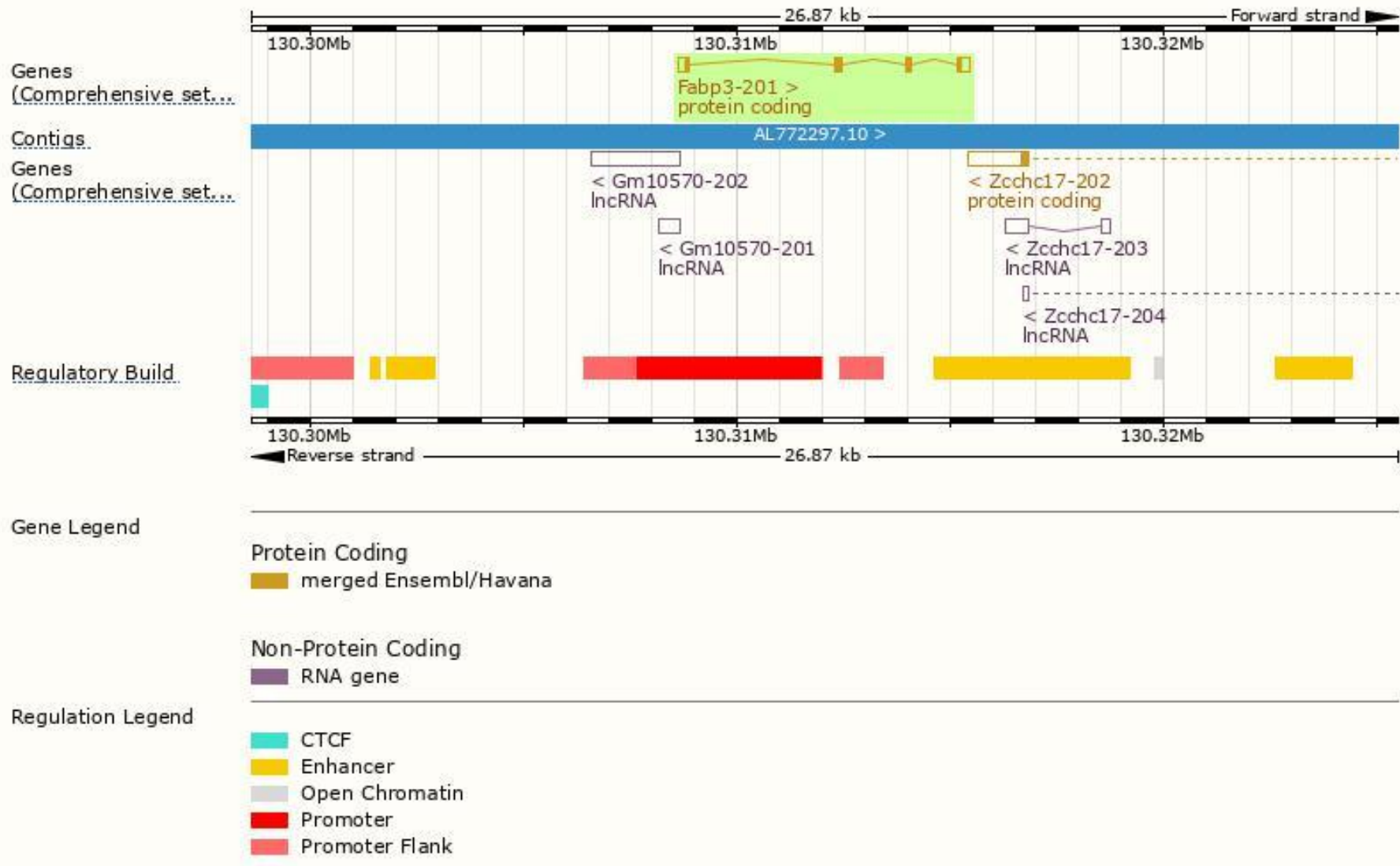
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fabp3-201	ENSMUST00000070532.7	823	133aa	Protein coding	CCDS18709	P11404 Q5EBJ0	TSL:1 GENCODE basic APPRIS P1

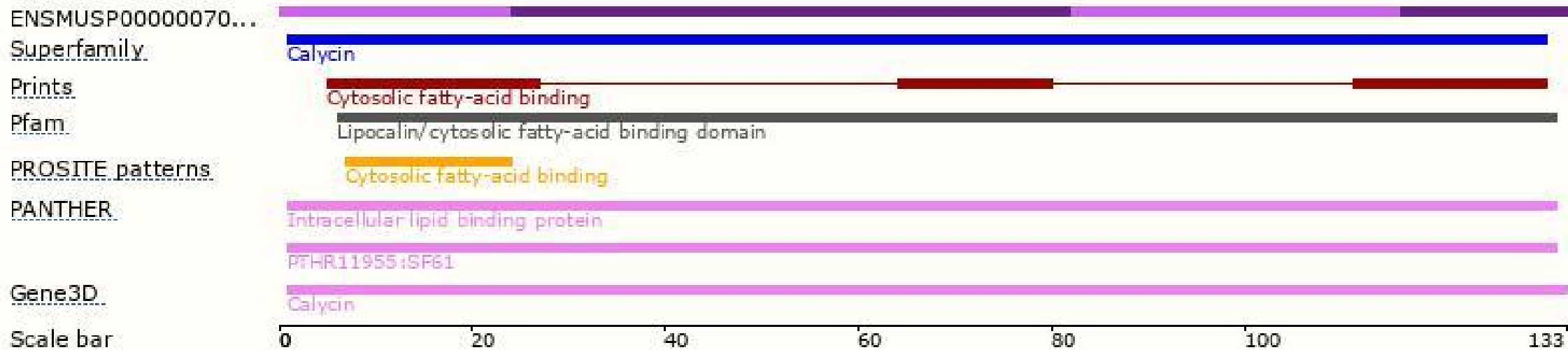
The strategy is based on the design of *Fabp3-201* transcript, The transcription is shown below



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, Inactivation of this locus results in impaired fatty acid utilization. Homozygous null mice show exercise intolerance and cardiac hypertrophy.

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

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