

Atf6 Cas9-KO Strategy

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Project Overview

Project Name

Atf6

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Atf6* gene. The schematic diagram is as follows:



- The *Atf6* gene has 2 transcripts. According to the structure of *Atf6* gene, exon2 of *Atf6-201* (ENSMUST00000027974.6) transcript is recommended as the knockout region. The region contains 77bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Atf6* gene. The brief process is as follows: CRISPR/Cas9 system w

- According to the existing MGI data, Mice homozygous for a null allele exhibit increased sensitivity to dithiothreitol, thapsigargin, and tunicamycin. Mice homozygous for a conditional allele activated in islet cells exhibit reduced sensitivity to TUDCA.
- The *Atf6* gene is located on the Chr1. 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)

Atf6 activating transcription factor 6 [Mus musculus (house mouse)]

Gene ID: 226641, updated on 26-Mar-2019

Summary



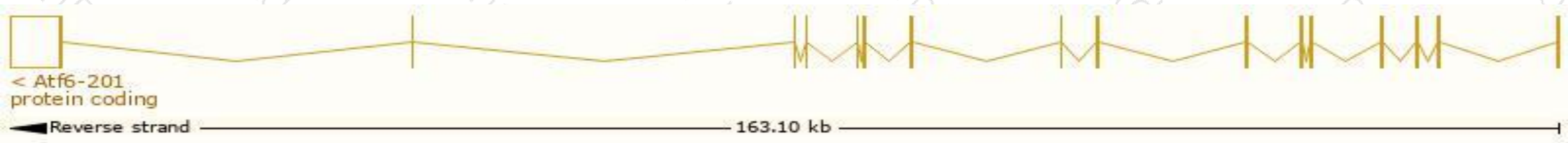
Official Symbol	Atf6 provided by MGI
Official Full Name	activating transcription factor 6 provided by MGI
Primary source	MGI:MGI:1926157
See related	Ensembl:ENSMUSG00000026663
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	9130025P16Rik, 9630036G24, AA789574, Atf6alpha, ESTM49
Expression	Ubiquitous expression in heart adult (RPKM 10.2), genital fat pad adult (RPKM 8.5) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

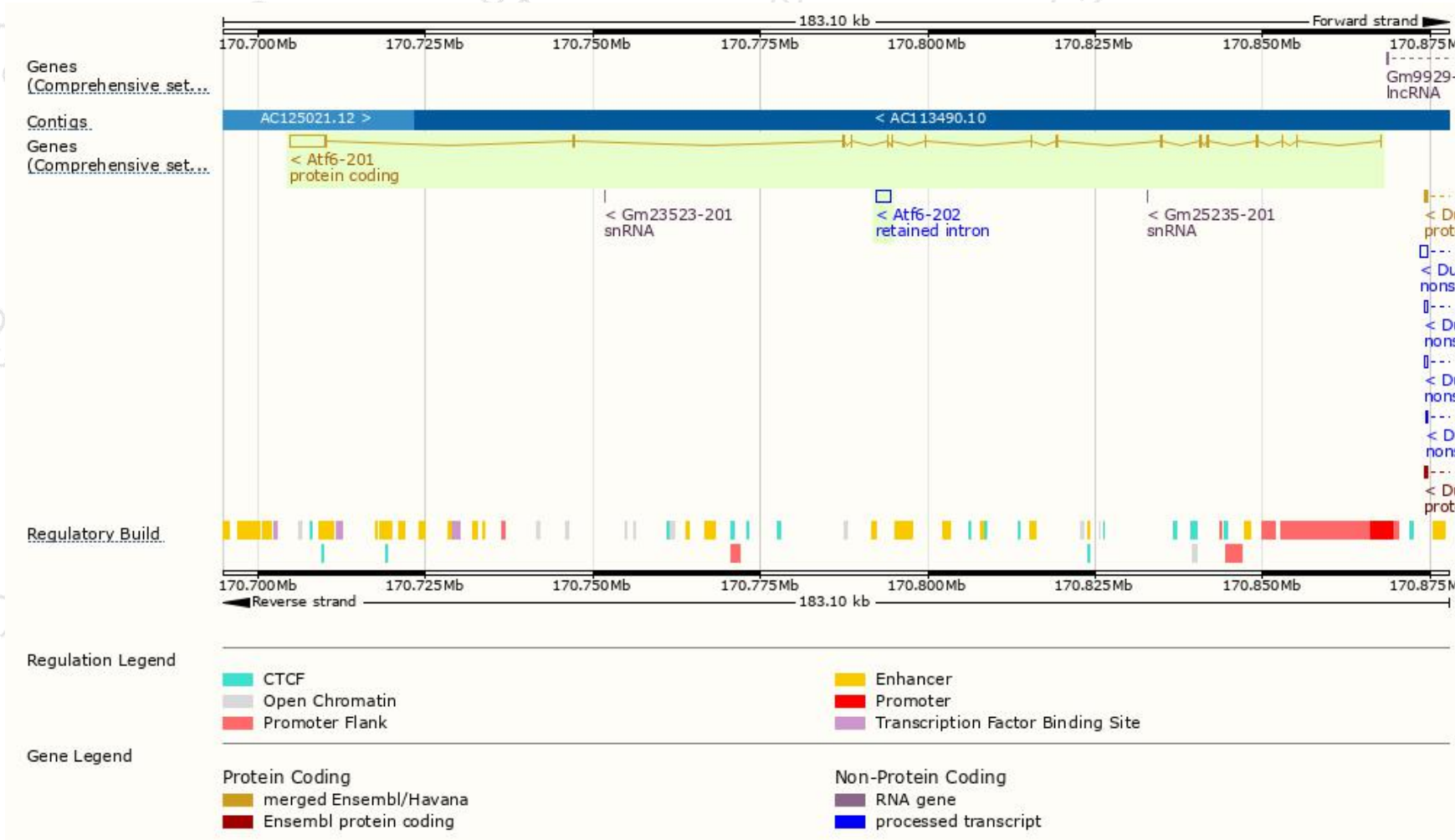
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Atf6-201	ENSMUST00000027974.6	7246	656aa	Protein coding	CCDS56653	F6VAN0	TSL:1 GENCODE basic APPRIS P1
Atf6-202	ENSMUST00000182787.1	2085	No protein	Retained intron	-	-	TSL:NA

The strategy is based on the design of *Atf6-201* transcript,The transcription is shown below



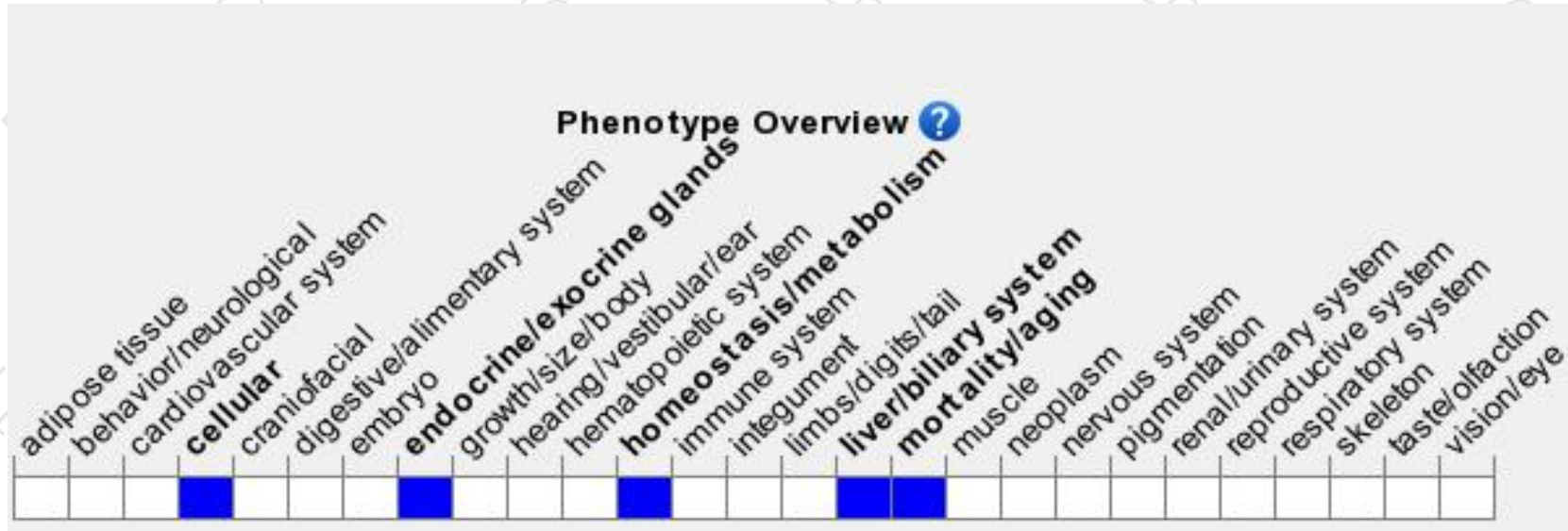
Genomic location distribution



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 null allele exhibit increased sensitivity to dithiothreitol, thapsigargin, and tunicamycin. Mice homozygous for a conditional allele activated in islet cells exhibit reduced sensitivity to TUDCA.

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

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