

Sntb1 Cas9-CKO Strategy

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Project Overview

Project Name

Sntb1

Project type

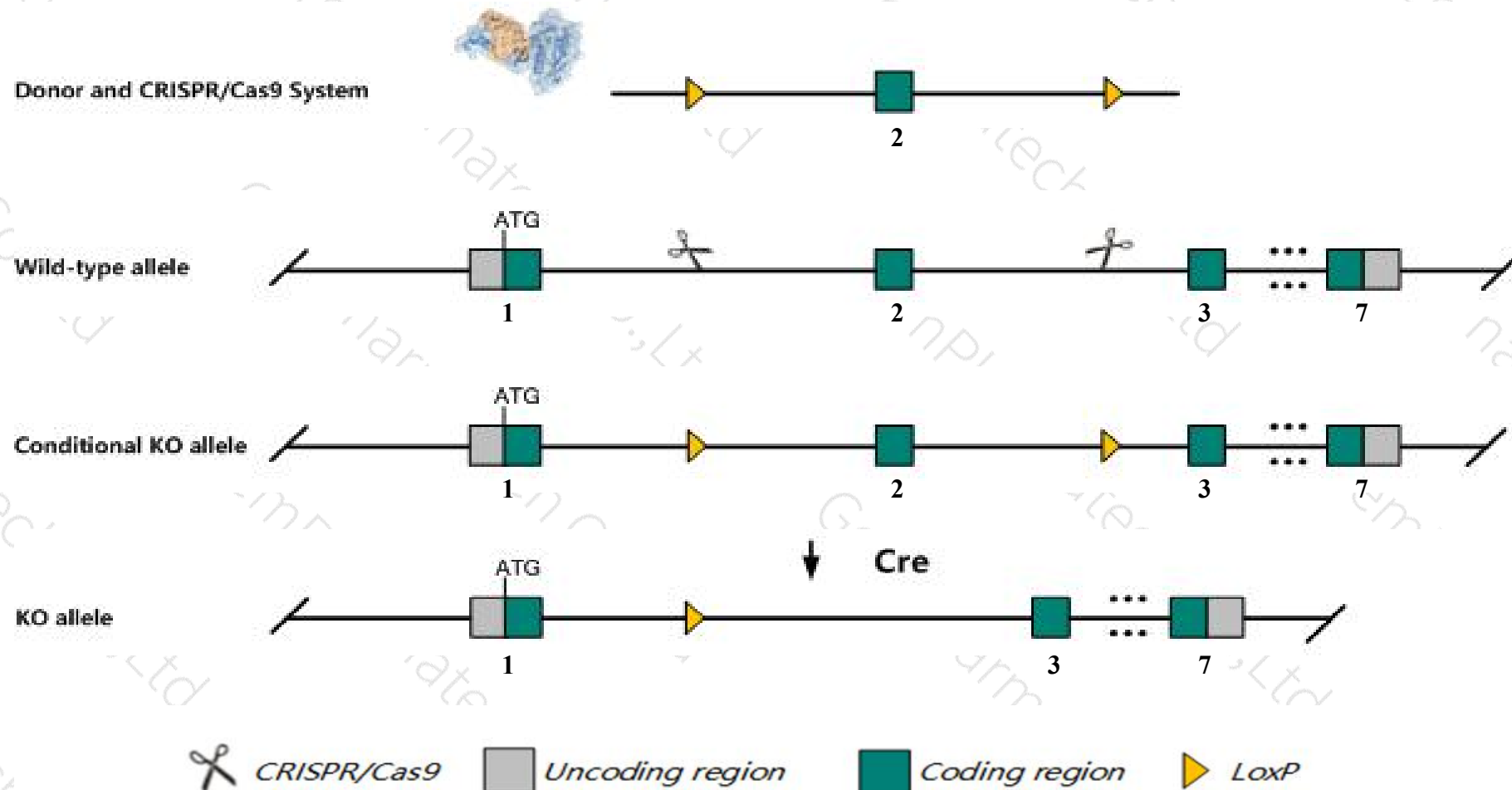
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Sntb1* gene. The schematic diagram is as follows:



Technical routes

- The *Sntb1* gene has 3 transcripts. According to the structure of *Sntb1* gene, exon2 of *Sntb1-201* (ENSMUST00000039769.12) transcript is recommended as the knockout region. The region contains 217bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Sntb1* 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 *Sntb1* gene is located on the Chr15. 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.
- Some amino acids will remain at the N-terminus and some functions may be retained.
- 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)

Sntb1 syntrophin, basic 1 [Mus musculus (house mouse)]

Gene ID: 20649, updated on 31-Jan-2019

Summary



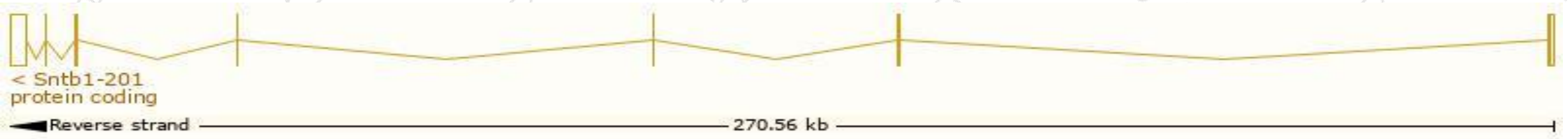
Official Symbol	Sntb1 provided by MGI
Official Full Name	syntrophin, basic 1 provided by MGI
Primary source	MGI:MGI:101781
See related	Ensembl:ENSMUSG00000060429
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Broad expression in subcutaneous fat pad adult (RPKM 7.5), adrenal adult (RPKM 7.5) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

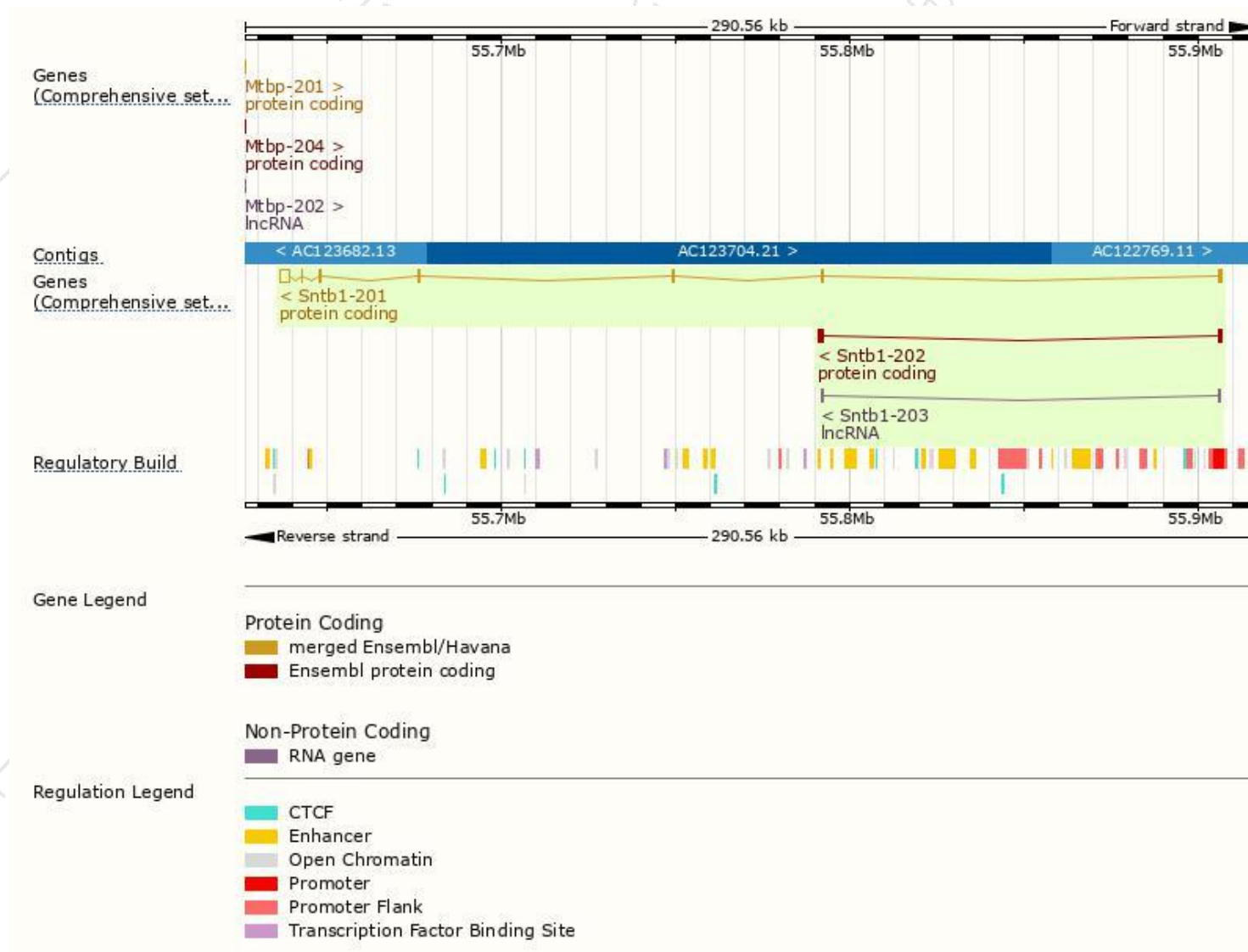
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Sntb1-201	ENSMUST00000039769.12	4919	537aa	Protein coding	CCDS27479	Q99L88	TSL:1 GENCODE basic APPRIS P1
Sntb1-202	ENSMUST00000110200.2	1225	276aa	Protein coding	-	Q99L88	TSL:1 GENCODE basic
Sntb1-203	ENSMUST00000140574.1	600	No protein	lncRNA	-	-	TSL:2

The strategy is based on the design of *Sntb1-201* 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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