

Sntb2 Cas9-KO Strategy

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Design Date:

2020-2-14

Project Overview

Project Name

Sntb2

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Sntb2* gene. The schematic diagram is as follows:



- The *Sntb2* gene has 3 transcripts. According to the structure of *Sntb2* gene, exon2 of *Sntb2-203* (ENSMUST00000212524.1) transcript is recommended as the knockout region. The region contains 214bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Sntb2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous null mice have no overt phenotype. They are fertile and motile with no signs of muscular dystrophy.
- The *Sntb2* gene is located on the Chr8. 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)

Sntb2 syntrophin, basic 2 [Mus musculus (house mouse)]

Gene ID: 20650, updated on 6-Feb-2019

Summary



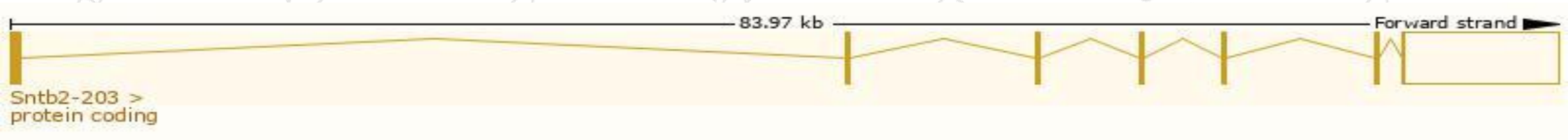
Official Symbol	Sntb2 provided by MGI
Official Full Name	syntrophin, basic 2 provided by MGI
Primary source	MGI:MGI:101771
See related	Ensembl:ENSMUSG00000041308
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	SNT3, SNTL, Snt2
Expression	Broad expression in ovary adult (RPKM 9.9), testis adult (RPKM 7.1) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

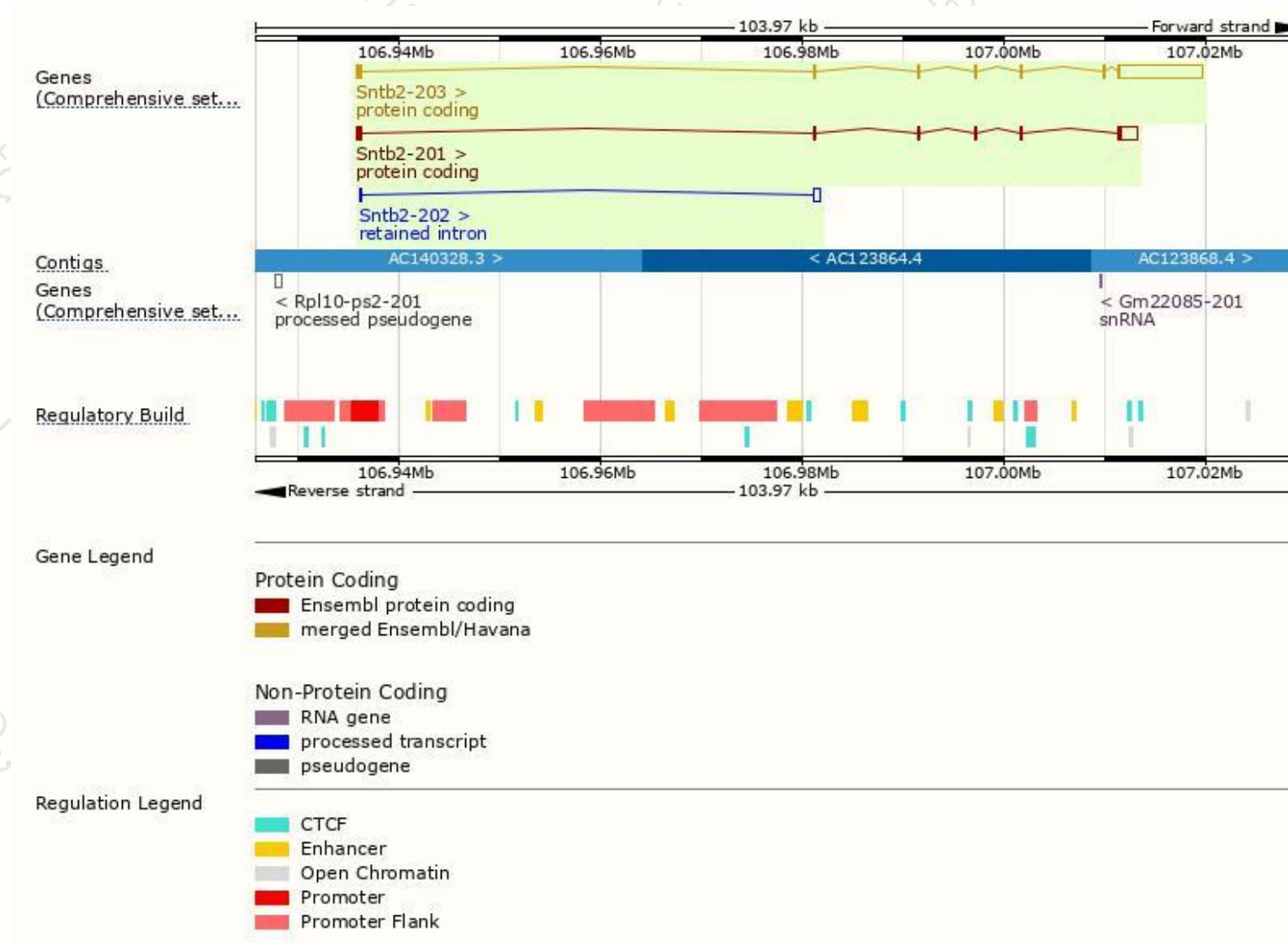
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Sntb2-203	ENSMUST00000212524.1	9945	520aa	Protein coding	CCDS22642	Q542S9 Q61235	TSL:1 GENCODE basic APPRIS P2
Sntb2-201	ENSMUST00000047425.4	3225	531aa	Protein coding	-	B7ZNU9	TSL:1 GENCODE basic APPRIS ALT2
Sntb2-202	ENSMUST00000212298.1	792	No protein	Retained intron	-	-	TSL:2

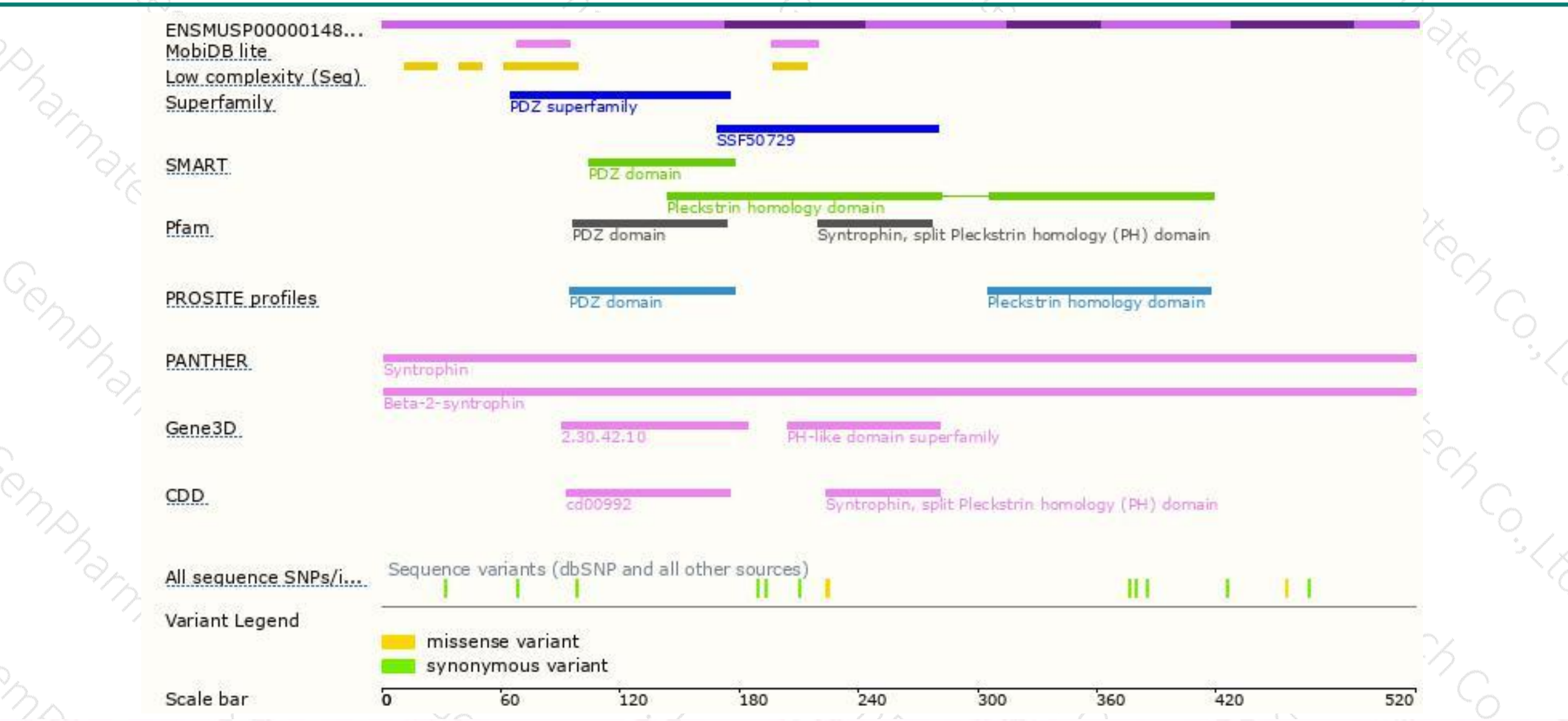
The strategy is based on the design of *Sntb2-203* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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