

$Snx31$ Cas9-KO Strategy

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Project Overview

Project Name

$Snx31$

Project type

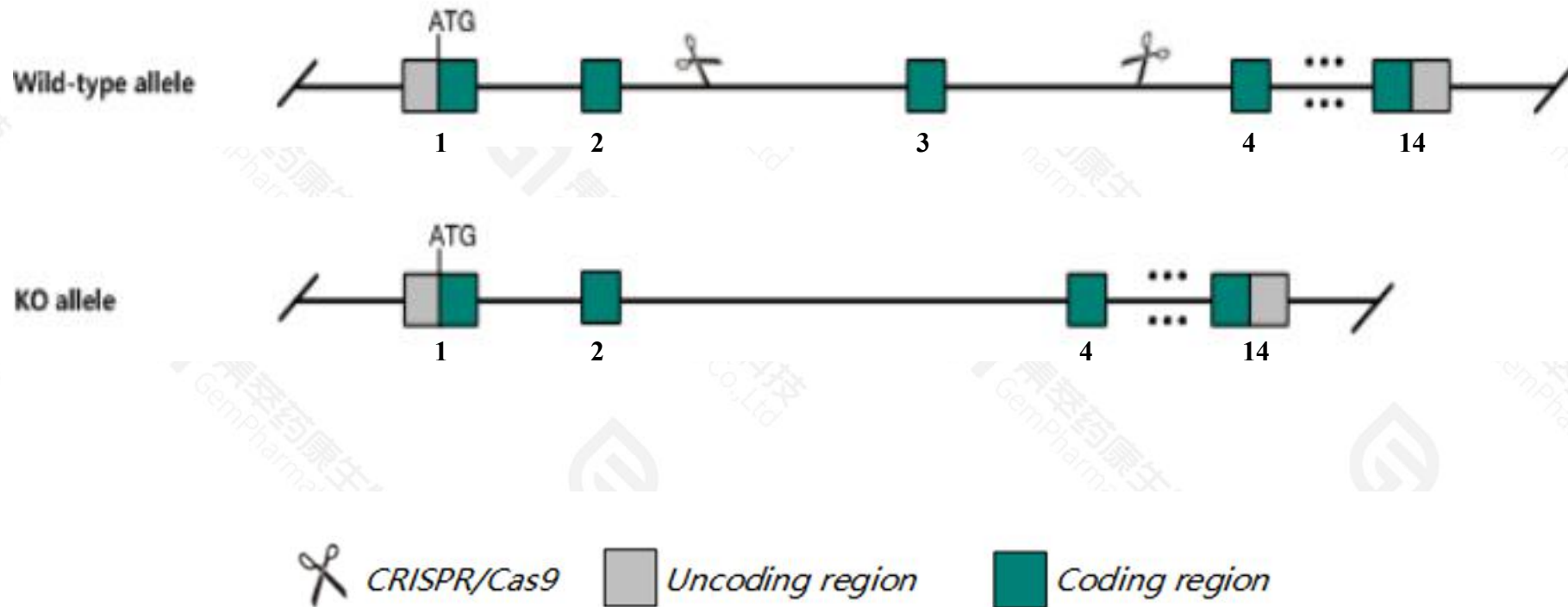
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Snx31* gene. The schematic diagram is as follows:



- The *Snx31* gene has 4 transcripts. According to the structure of *Snx31* gene, exon3 of *Snx31*-203(ENSMUST00000161202.8) transcript is recommended as the knockout region. The region contains 115bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Snx31* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice

- The *Snx31* 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.
- 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)

Snx31 sorting nexin 31 [Mus musculus (house mouse)]

Gene ID: 66696, updated on 17-Dec-2020

Summary



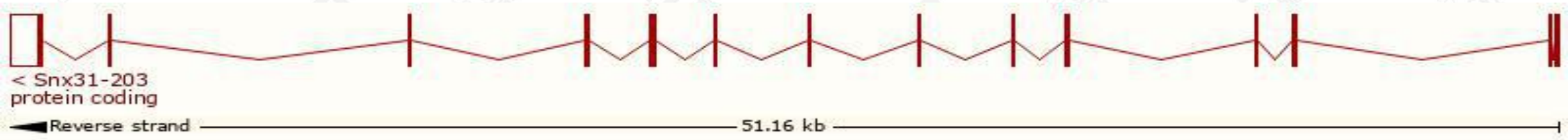
Official Symbol	Snx31 provided by MGI
Official Full Name	sorting nexin 31 provided by MGI
Primary source	MGI:MGI:1913946
See related	Ensembl:ENSMUSG00000013611
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4631426E05Rik
Expression	Restricted expression toward bladder adult (RPKM 80.8) See more
Orthologs	human all

Transcript information (Ensembl)

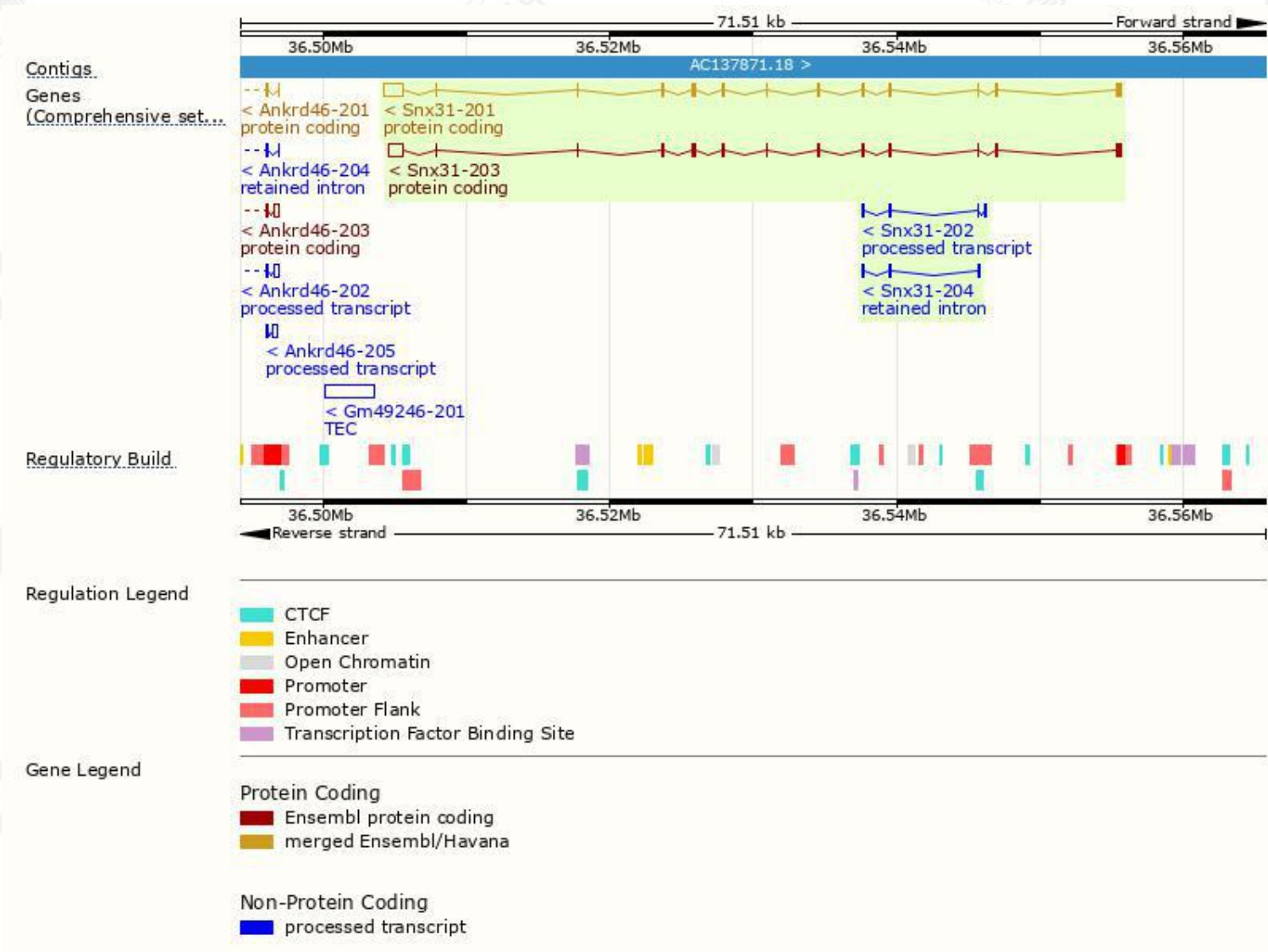
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Snx31-201	ENSMUST00000013755.12	2696	438aa	Protein coding	CCDS27430		TSL:1 , GENCODE basic , APPRIS P3 ,
Snx31-203	ENSMUST000000161202.8	2344	439aa	Protein coding	CCDS84166		TSL:1 , GENCODE basic , APPRIS ALT2 ,
Snx31-202	ENSMUST000000160936.3	430	No protein	Processed transcript	-		TSL:3 ,
Snx31-204	ENSMUST000000162888.2	340	No protein	Retained intron	-		TSL:3 ,

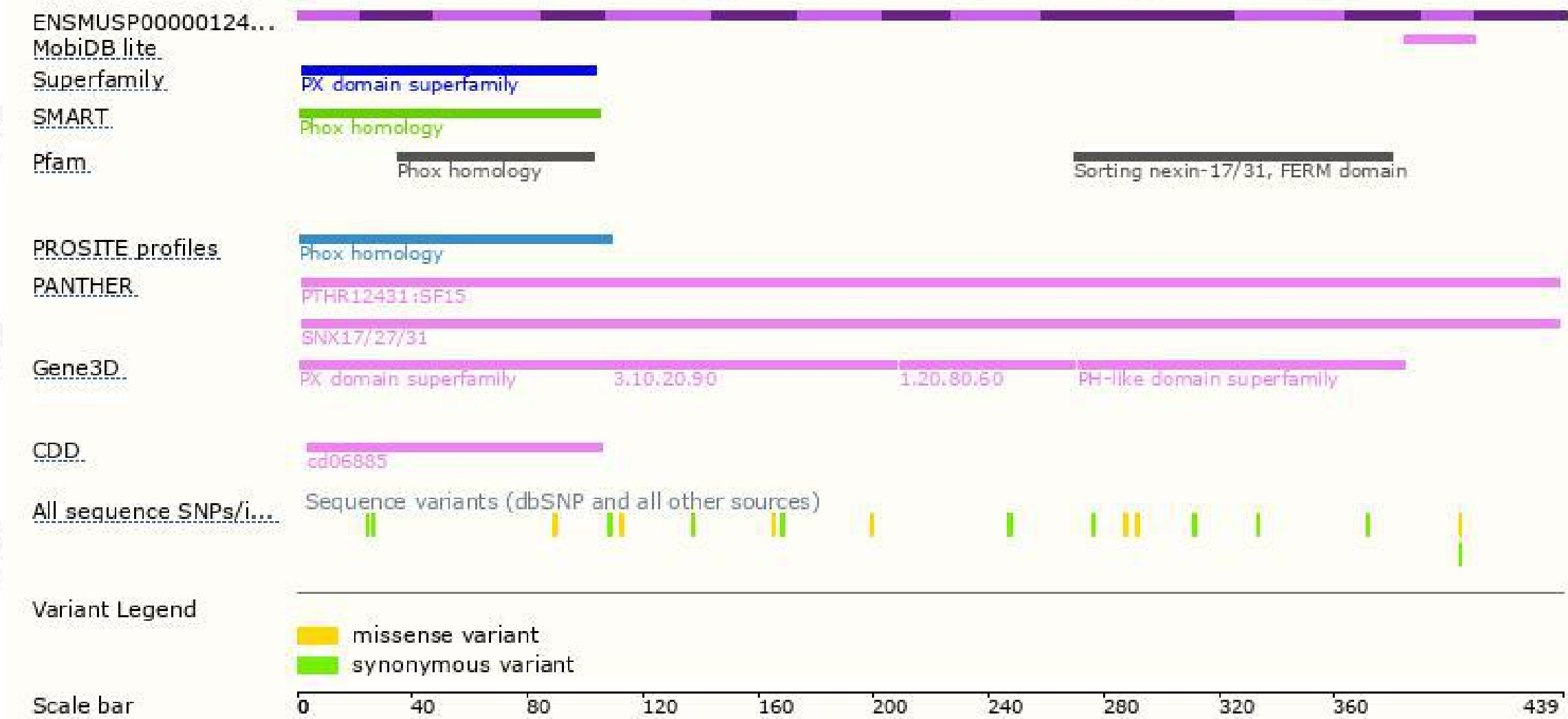
The strategy is based on the design of *Snx31-203* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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