

***Zbtb6* Cas9-KO Strategy**

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Project Overview

Project Name

Zbtb6

Project type

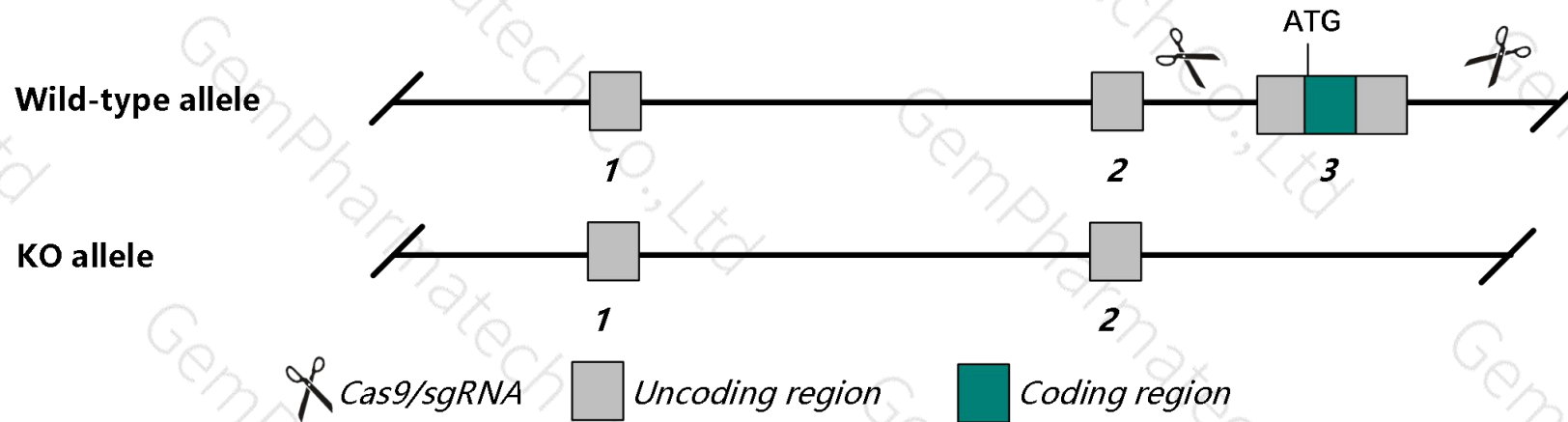
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zbtb6* gene. The schematic diagram is as follows:



- The *Zbtb6* gene has 2 transcripts. According to the structure of *Zbtb6* gene, exon3 of *Zbtb6*-202(ENSMUST00000112932.1) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zbtb6* 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.

- The *Zbtb6* gene is located on the Chr2. 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.
- The knockout region is near to the N-terminal of *Rc3h2* gene and the C-terminal of *Zbtb6* gene, this strategy may influence the regulatory function of the N-terminal of *Rc3h2* gene and the C-terminal of *Zbtb6* gene.
- 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)

Zbtb6 zinc finger and BTB domain containing 6 [Mus musculus (house mouse)]

Gene ID: 241322, updated on 13-Mar-2020

Summary



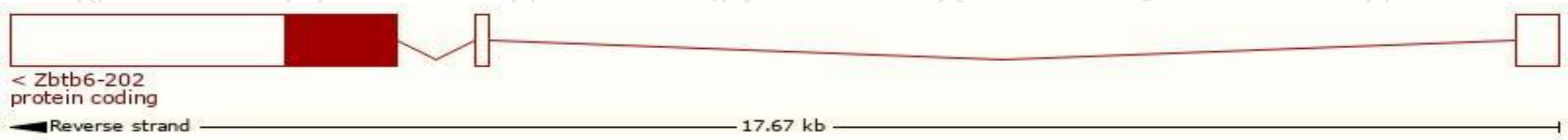
Official Symbol	Zbtb6 provided by MGI
Official Full Name	zinc finger and BTB domain containing 6 provided by MGI
Primary source	MGI:MGI:2442998
See related	Ensembl:ENSMUSG00000066798
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	A830092L04Rik, ZID, Zfp482
Expression	Broad expression in CNS E18 (RPKM 5.5), whole brain E14.5 (RPKM 4.7) and 24 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

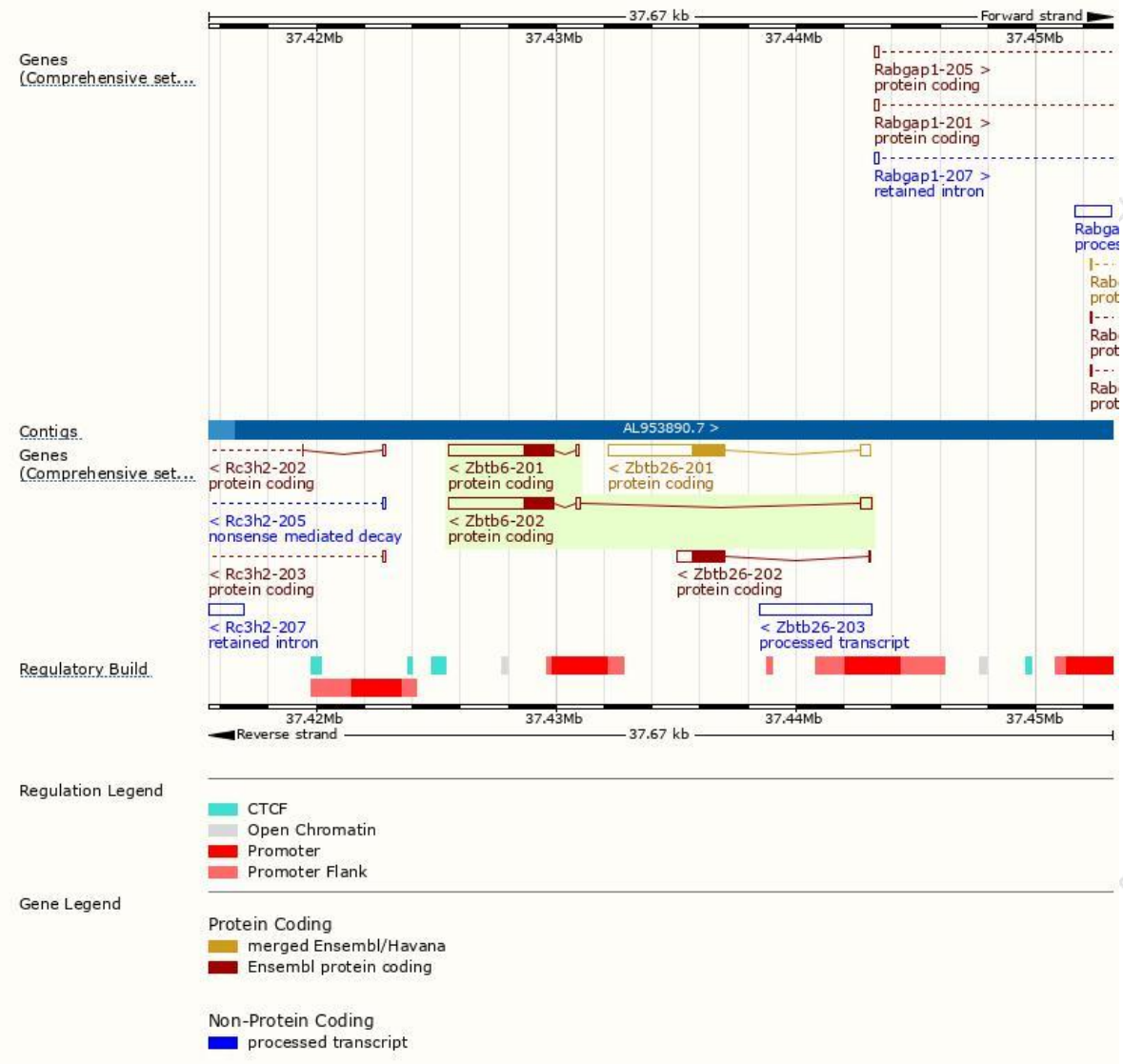
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zbtb6-202	ENSMUST00000112932.1	5077	423aa	Protein coding	CCDS16000	Q8K088	TSL:2 GENCODE basic APPRIS P1
Zbtb6-201	ENSMUST00000053098.5	4535	423aa	Protein coding	CCDS16000	Q8K088	TSL:1 GENCODE basic APPRIS P1

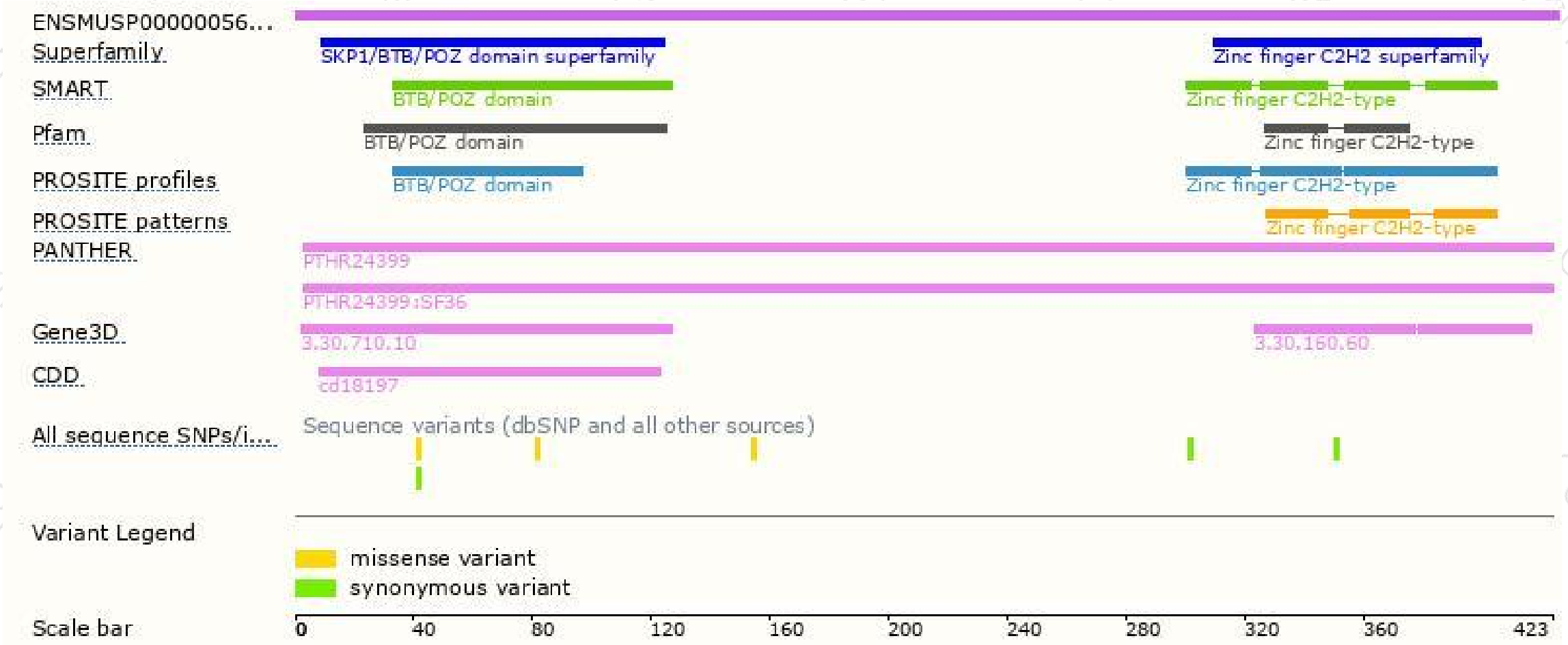
The strategy is based on the design of *Zbtb6-202* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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