

Zbtb41 Cas9-KO Strategy

Designer: Xueting Zhang

Reviewer: Daohua Xu

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Project Overview



Project Name

Zbtb41

Project type

Cas9-KO

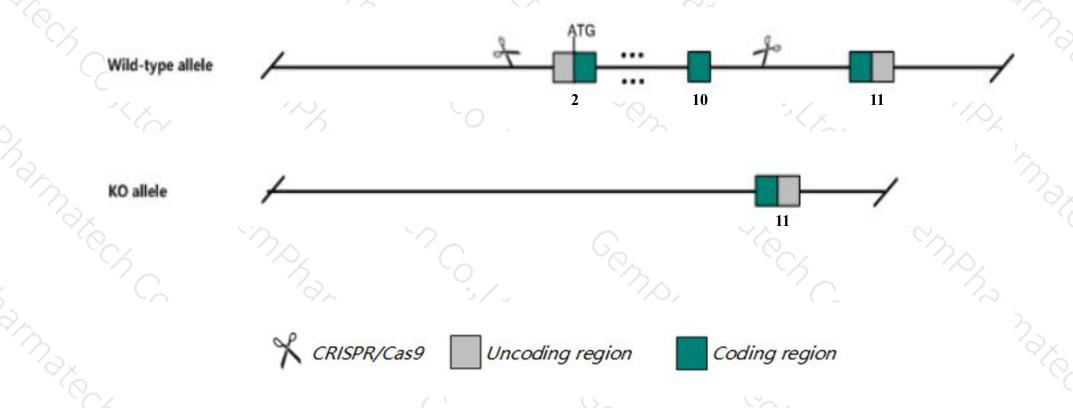
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Zbtb41 gene. The schematic diagram is as follows:



Technical routes



- ➤ The Zbtb41 gene has 3 transcripts. According to the structure of Zbtb41 gene, exon2-exon10 of Zbtb41-201(ENSMUST00000039867.9) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zbtb41* 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 Zbtb41 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)



Zbtb41 zinc finger and BTB domain containing 41 [Mus musculus (house mouse)]

Gene ID: 226470, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Zbtb41 provided by MGI

Official Full Name zinc finger and BTB domain containing 41 provided by MGI

Primary source MGI:MGI:2444487

See related Ensembl: ENSMUSG00000033964

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 8430415N23Rik, 9430031N01, 9830132G07Rik, Al316857

Expression Ubiquitous expression in CNS E18 (RPKM 5.4), whole brain E14.5 (RPKM 5.0) and 25 other tissuesSee more

Orthologs <u>human all</u>

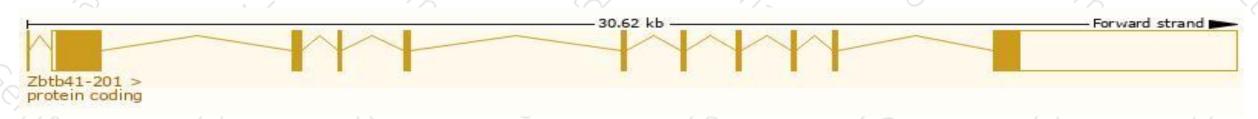
Transcript information (Ensembl)



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

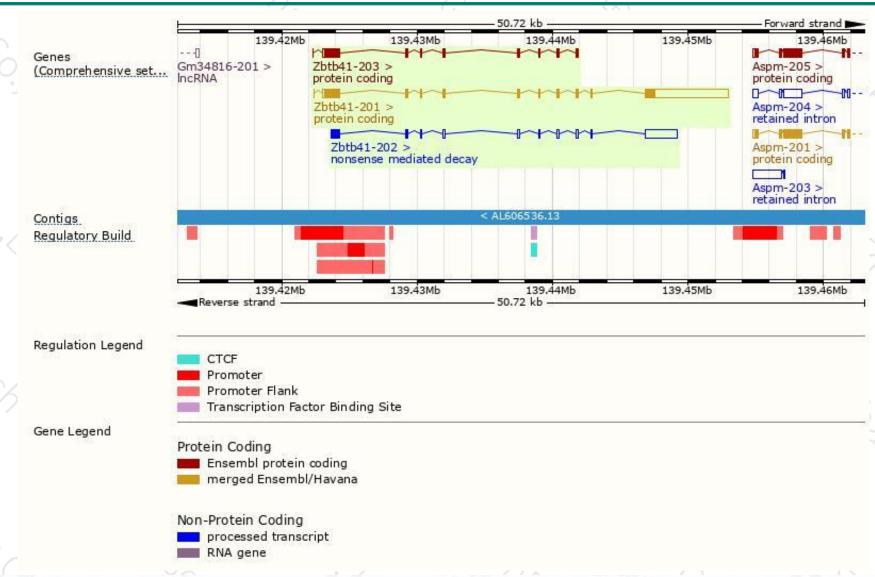
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
200000000000000000000000000000000000000	ENSMUST00000039867.9	8361	908aa	Protein coding	CCDS48385	Q811F1	TSL:1 GENCODE basic APPRIS P1
	ENSMUST00000200243.4		657aa	Protein coding	-	A0A0G2JEJ7	CDS 3' incomplete TSL:5
Zbtb41-202	ENSMUST00000199011.1	3803	219aa	Nonsense mediated decay	2	A0A0G2JH01	CDS 5' incomplete TSL:1

The strategy is based on the design of Zbtb41-201 transcript, the transcription is shown below:



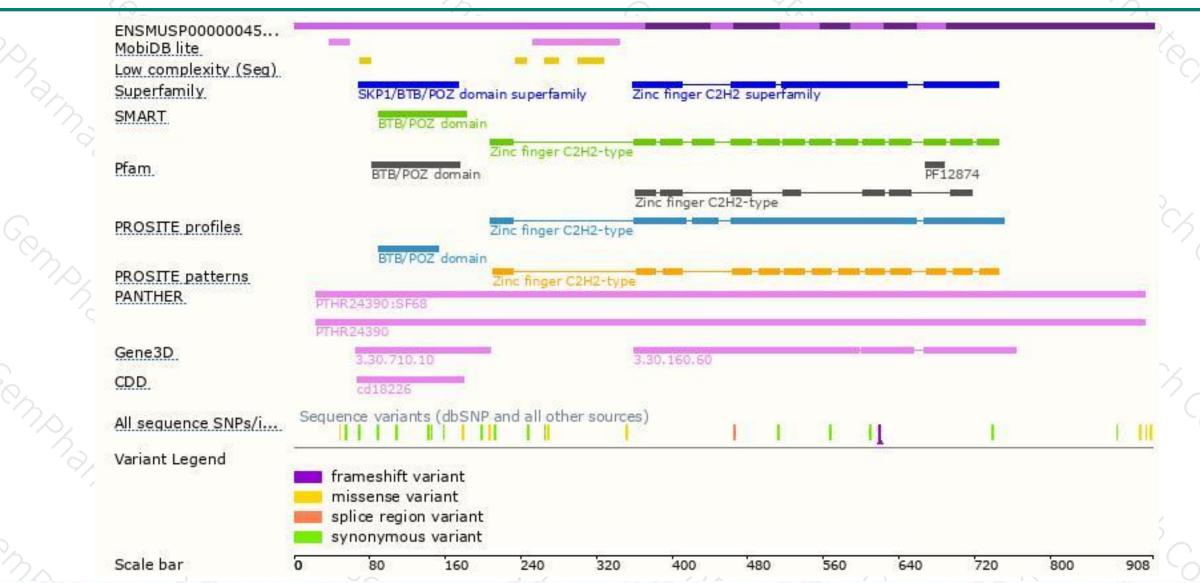
Genomic location distribution





Protein domain







If you have any questions, you are welcome to inquire. Tel: 400-9660890





