

Zbtb48 Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview



Project Name

Zbtb48

Project type

Cas9-CKO

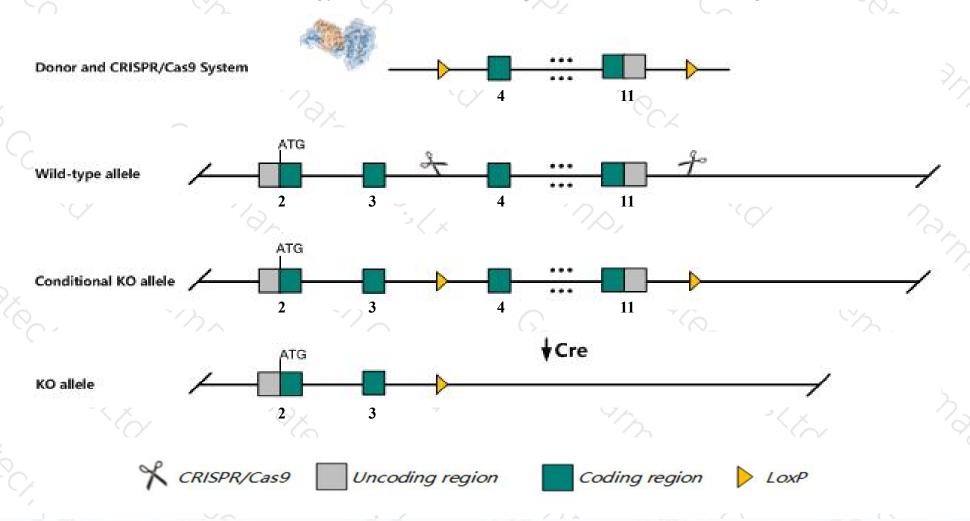
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Zbtb48* gene. The schematic diagram is as follows:



Technical routes



- The *Zbtb48* gene has 8 transcripts. According to the structure of *Zbtb48* gene, exon4-exon11 of *Zbtb48-201* (ENSMUST00000066715.10) transcript is recommended as the knockout region. The region contains 1135bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zbtb48* 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 Zbtb48 gene is located on the Chr4. 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 n-terminal amino acid residues are about half of the total, and the function of the protein may be preserved.
- > 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)



Zbtb48 zinc finger and BTB domain containing 48 [Mus musculus (house mouse)]

Gene ID: 100090, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Zbtb48 provided by MGI

Official Full Name zinc finger and BTB domain containing 48 provided by MGI

Primary source MGI:MGI:2140248

See related Ensembl:ENSMUSG00000028952

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 0610011D15Rik, Al327031, Hkr3, TZAP

Expression Ubiquitous expression in thymus adult (RPKM 8.7), limb E14.5 (RPKM 7.7) and 28 other tissuesSee more

Orthologs <u>human</u> all

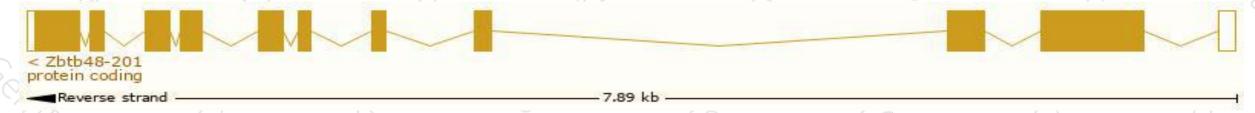
Transcript information (Ensembl)



The gene has 8 transcripts, all transcripts are shown below:

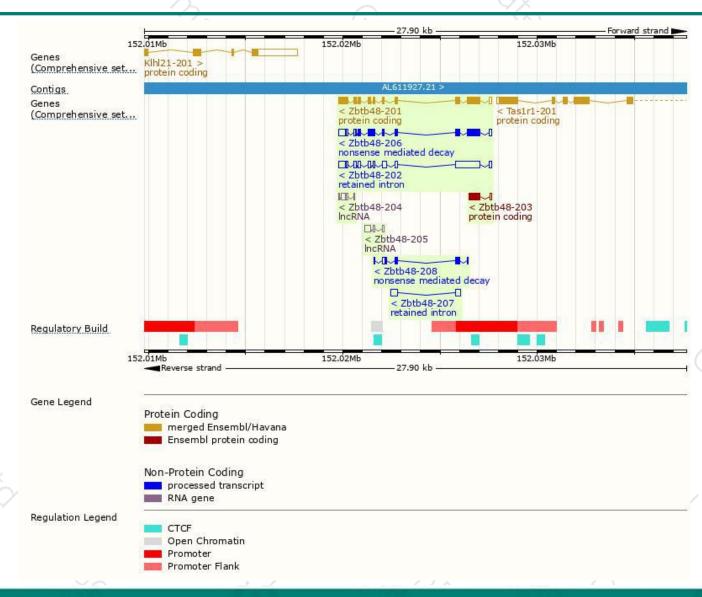
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zbtb48-201	ENSMUST00000066715.10	2218	<u>681aa</u>	Protein coding	CCDS38982	Q1H9T6	TSL:1 GENCODE basic APPRIS P1
Zbtb48-203	ENSMUST00000131935.1	655	<u>193aa</u>	Protein coding		A0A0A0MQG8	CDS 3' incomplete TSL:3
Zbtb48-206	ENSMUST00000155389.7	2312	<u>547aa</u>	Nonsense mediated decay	(20)	Q1H9T6	TSL:2
Zbtb48-208	ENSMUST00000156748.1	656	<u>164aa</u>	Nonsense mediated decay	120	F6ZA56	CDS 5' incomplete TSL:3
Zbtb48-202	ENSMUST00000123696.7	2709	No protein	Retained intron	153	35	TSL:1
Zbtb48-207	ENSMUST00000155441.1	592	No protein	Retained intron		195	TSL:3
Zbtb48-205	ENSMUST00000147895.1	502	No protein	IncRNA	(20)	<u> </u>	TSL:3
Zbtb48-204	ENSMUST00000136212.1	406	No protein	IncRNA	150	62	TSL:3

The strategy is based on the design of Zbtb48-201 transcript, The transcription is shown below



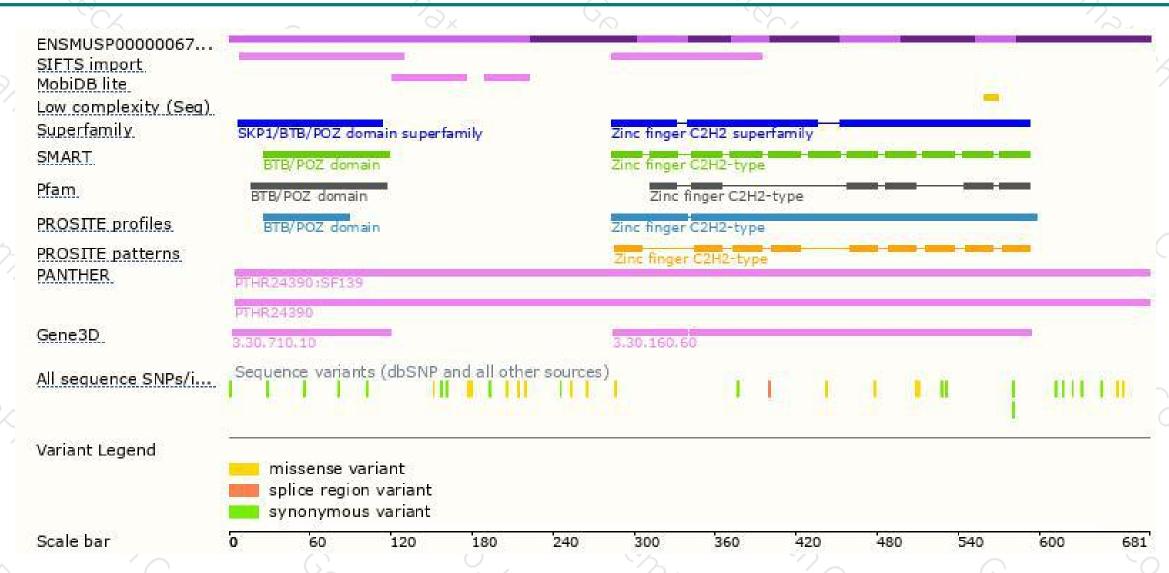
Genomic location distribution





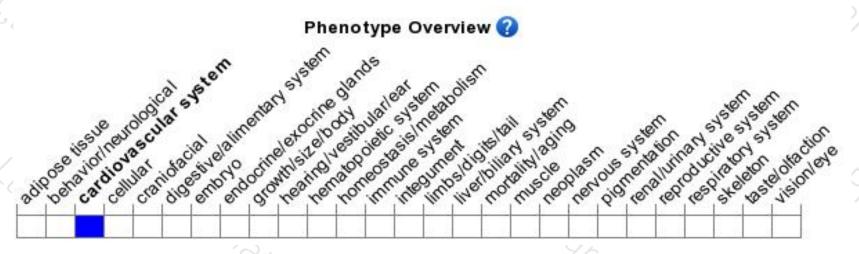
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).



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





