

Tekt5 Cas9-CKO Strategy

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Project Overview



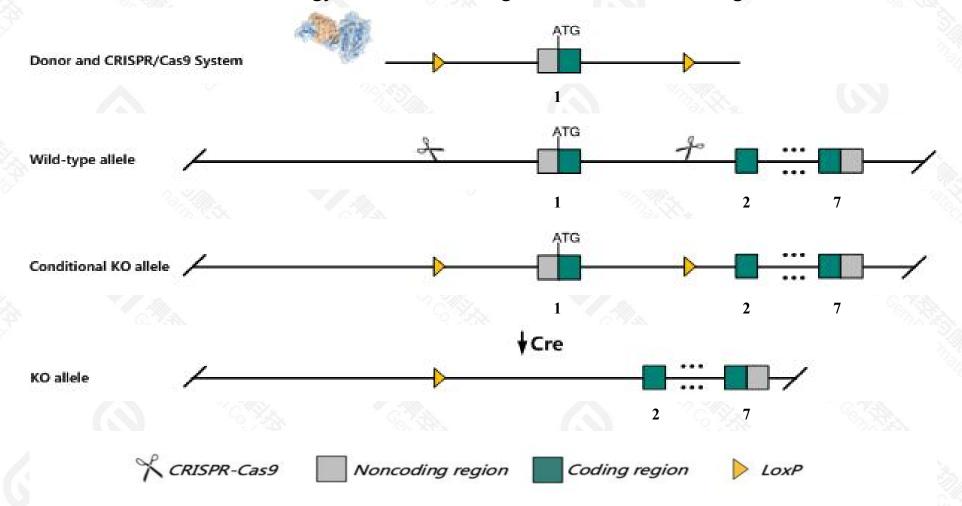
Project Name	Tekt5			
Project type	Cas9-CKO			
Strain background	C57BL/6JGpt			

GemPharmatech Co., Ltd.

Conditional Knockout strategy



This model will use CRISPR-Cas9 technology to edit the *Tekt5* gene. The schematic diagram is as follows:



Technical routes



- The *Tekt5* gene has 2 transcripts. According to the structure of *Tekt5* gene, exon1 of *Tekt5-201*(ENSMUST00000043415.13) 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 *Tekt5* 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 *Tekt5* gene is located on the Chr16. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Tekt5 tektin 5 [Mus musculus (house mouse)]

Gene ID: 70426, updated on 24-Apr-2022

Summary



Official Symbol Tekt5 provided by MGI
Official Full Name tektin 5 provided by MGI
Primary source MGI:MGI:1917676

See related Ensembl: ENSMUSG00000039179

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 3300001K11Rik

Expression Biased expression in testis adult (RPKM 92.6) and cerebellum adult (RPKM 4.3)See more

Orthologs <u>human all</u>

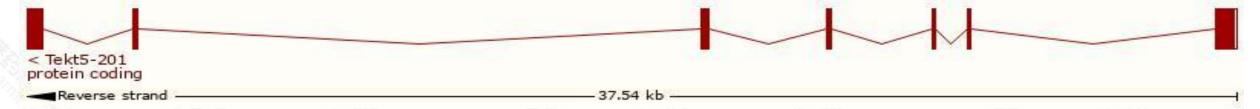
Transcript information (Ensembl)



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

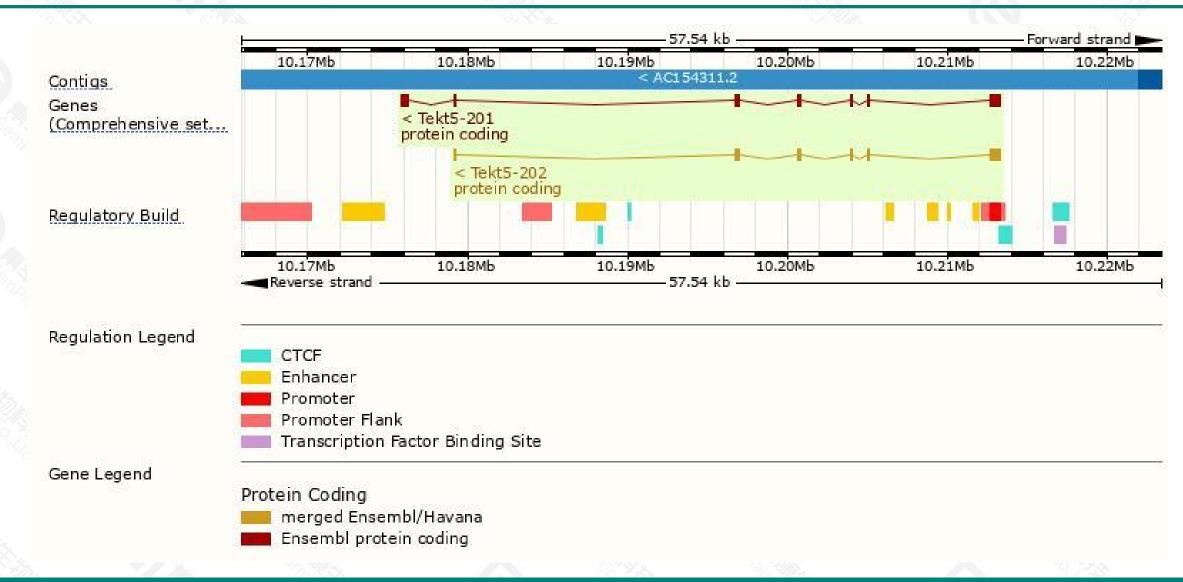
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tekt5-201	ENSMUST00000043415.13	1803	<u>557aa</u>	Protein coding	CCDS70683		TSL:1, GENCODE basic, APPRIS P1,
Tekt5-202	ENSMUST00000115831.2	1277	<u>415aa</u>	Protein coding	-		TSL:1, GENCODE basic,

The strategy is based on the design of *Tekt5-201* transcript, the transcription is shown below:



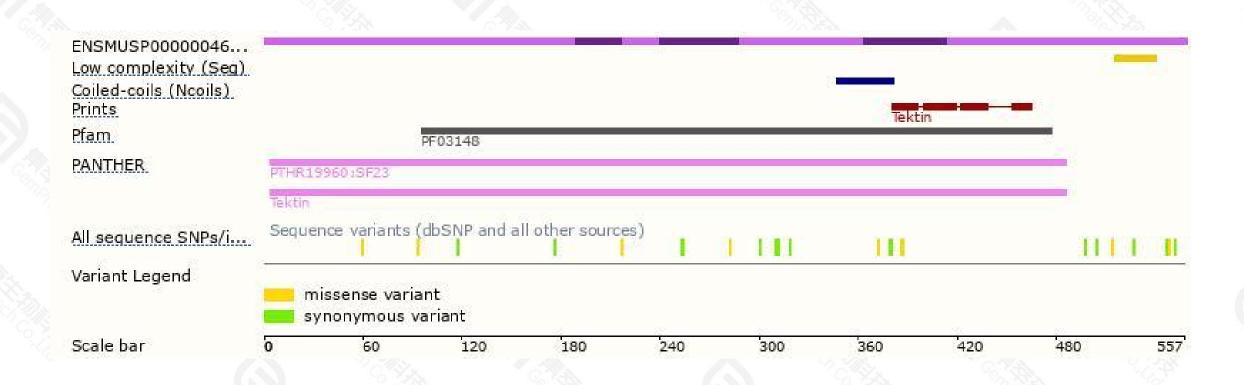
Genomic location distribution





Protein domain







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

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