

Tsen15 Cas9-KO Strategy

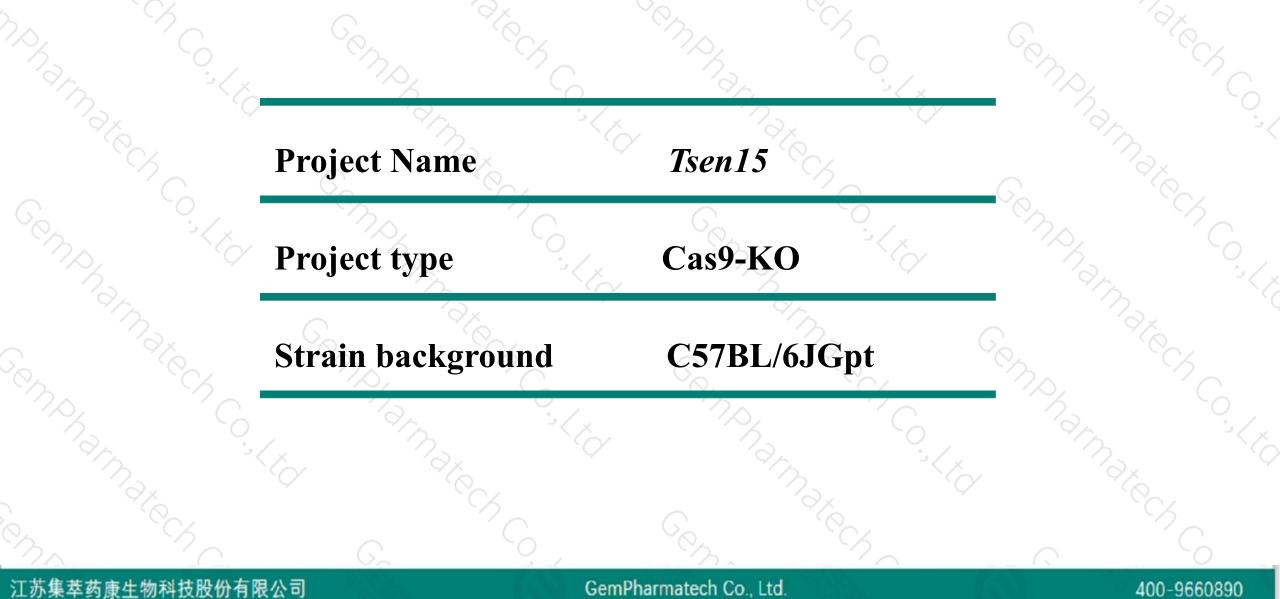
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Reviewer: Shanhong Tao

Design Date: 2021-3-29

Project Overview

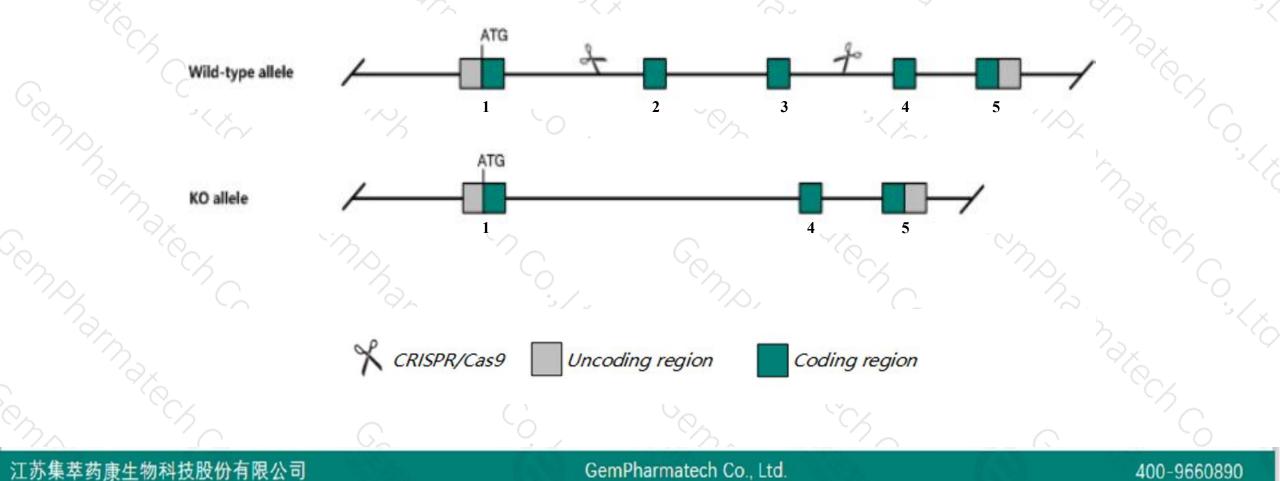




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Tsen15 gene. The schematic diagram is as follows:





➤ The *Tsen15* gene has 7 transcripts. According to the structure of *Tsen15* gene, exon2-exon3 of *Tsen15*-201(ENSMUST00000015124.14) transcript is recommended as the knockout region. The region contains 218bp coding sequence. Knock out the region will result in disruption of protein function.

➤ In this project we use CRISPR/Cas9 technology to modify *Tsen15* 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 *Tsen15* 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.

Notice

Gene information (NCBI)



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Tsen15 tRNA splicing endonuclease subunit 15 [Mus musculus (house mouse)]

Gene ID: 66637, updated on 13-Mar-2020

Summary

Constant of the	
Official Symbol	Tsen15 provided by MGI
Official Full Name	tRNA splicing endonuclease subunit 15 provided by MGI
Primary source	MGI:MGI:1913887
See related	Ensembl:ENSMUSG00000014980
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	5730449L18Rik, AL023077, Sen15
Expression	Ubiquitous expression in limb E14.5 (RPKM 14.7), placenta adult (RPKM 11.9) and 27 other tissues See more
Orthologs	human all

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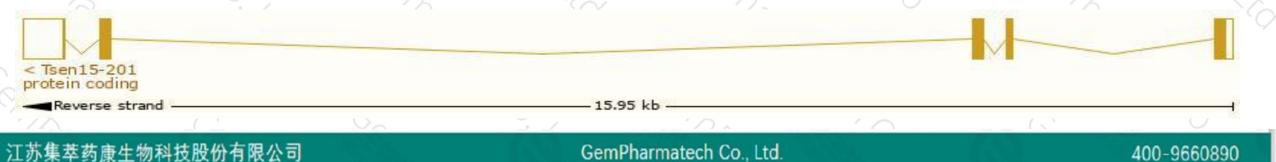
Transcript information (Ensembl)



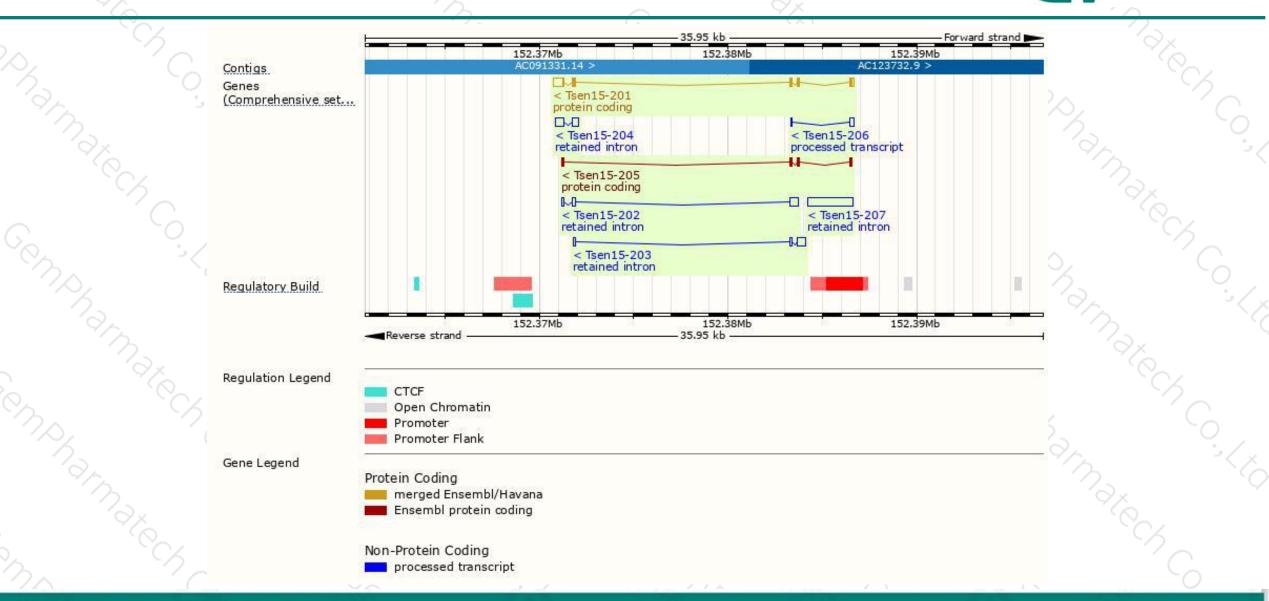
The gene has 7 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tsen15-201	ENSMUST00000015124.14	1133	<u>168aa</u>	Protein coding	CCDS15363	<u>G3X8S8</u>	TSL:1 GENCODE basic APPRIS P1
Tsen15-205	ENSMUST00000162371.1	431	<u>128aa</u>	Protein coding	, 8	E0CY96	TSL:5 GENCODE basic
Tsen15-206	ENSMUST00000163027.1	283	No protein	Processed transcript	10	23	TSL:3
Tsen15-207	ENSMUST00000188105.1	2418	No protein	Retained intron	<u>1</u> 0	20	TSL:NA
Tsen15-204	ENSMUST00000161717.1	775	No protein	Retained intron	54	-	TSL:2
Tsen15-202	ENSMUST00000159270.1	698	No protein	Retained intron	, 8	-	TSL:2
Tsen15-203	ENSMUST00000159717.1	651	No protein	Retained intron	29	20	TSL:5

The strategy is based on the design of *Tsen15-201* transcript, the transcription is shown below:



Genomic location distribution



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Protein domain



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	ENSMUSP00000015 MobiDB lite Superfamily	tRNA intron endonuclease, catalytic domain-like superfamily									
Pfam tRNA-splicing endonuclease subunit Sen15 PANTHER PTHR28582											
	Gene3D	PIN2030		tRNA endo	nuclease-like d	Iomain superfamil	Y.				
	All sequence SNPs/i	Sequence	variants (dbS	SNP and all ot	her sources)						
Gen.	Variant Legend		ense variant e region varia	int						30	
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If you have any questions, you are welcome to inquire. Tel: 400-9660890



