

Dnajb13 Cas9-CKO Strategy

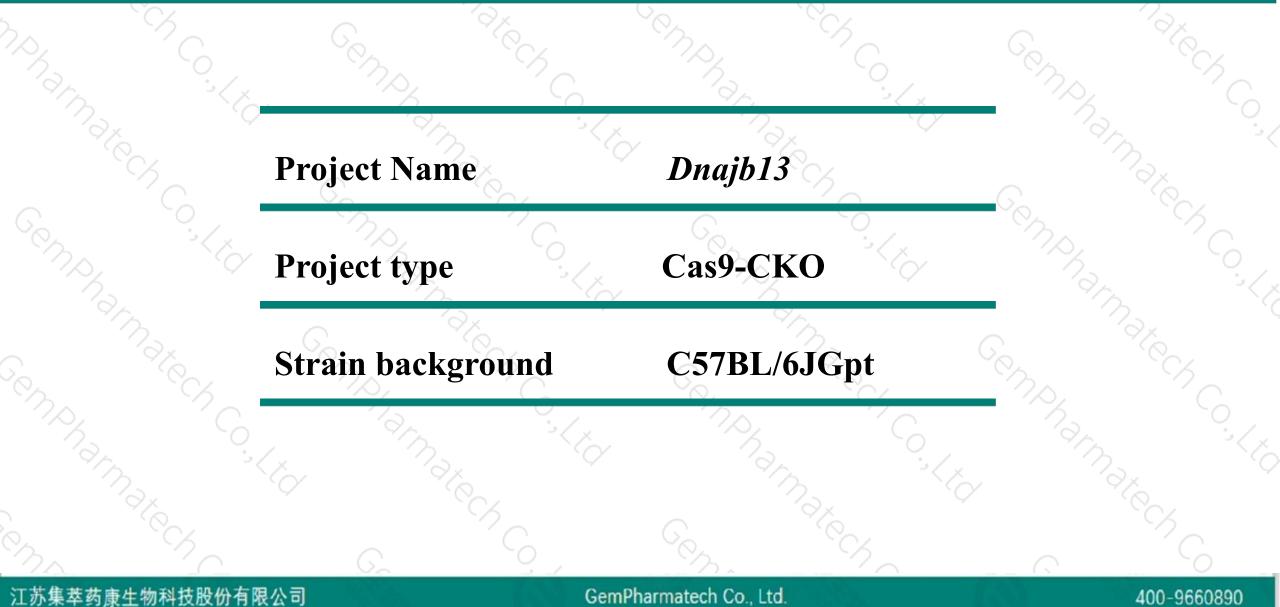
Designer: JiaYu

Reviewer: Xiaojing Li

Design Date: 2020-7-28

Project Overview



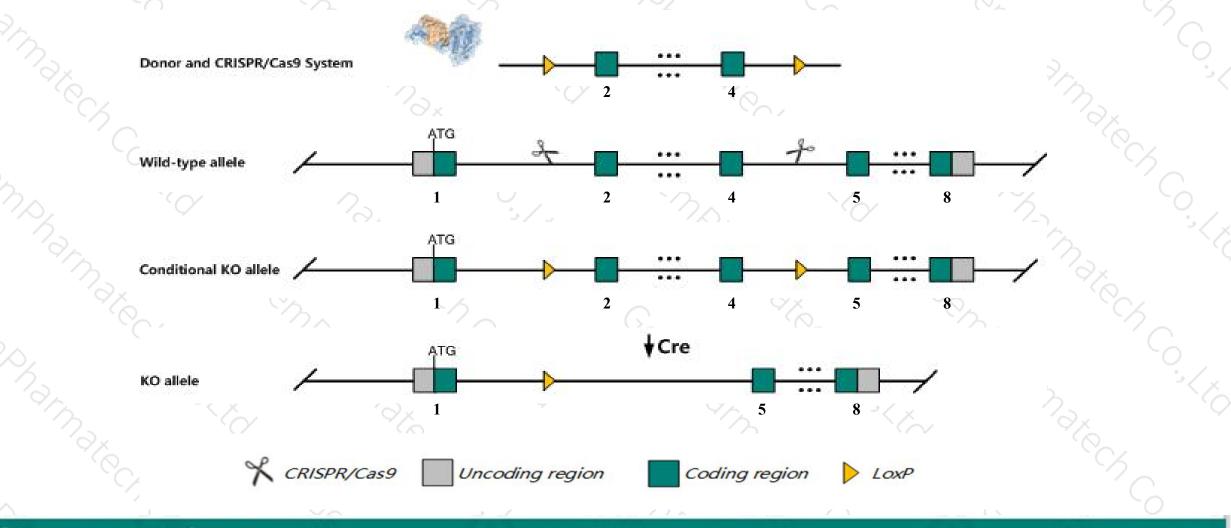


Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the *Dnajb13* gene. The schematic diagram is as follows:



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The Dnajb13 gene has 4 transcripts. According to the structure of Dnajb13 gene, exon2-exon4 of Dnajb13-204(ENSMUST00000207405.1) transcript is recommended as the knockout region. The region contains 424bp coding sequence. Knock out the region will result in disruption of protein function.

➤ In this project we use CRISPR/Cas9 technology to modify *Dnajb13* 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.



- > The *Dnajb13* gene is located on the Chr7. 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)



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Dnajb13 DnaJ heat shock protein family (Hsp40) member B13 [Mus musculus (house mouse)]

Gene ID: 69387, updated on 13-Mar-2020

Summary

Official Symbol	Dnajb13 provided by MGI
Official Full Name	DnaJ heat shock protein family (Hsp40) member B13 provided by MGI
Primary source	MGI:MGI:1916637
See related	Ensembl:ENSMUSG0000030708
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
	Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1700014P03Rik, Tsarg
Expression	Biased expression in testis adult (RPKM 160.4) and ovary adult (RPKM 6.8)See more
Orthologs	human all

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The gene has 4 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Dnajb13-204	ENSMUST00000207405.1	2592	<u>316aa</u>	Protein coding	CCDS21499	<u>Q80Y75</u>	TSL:1 GENCODE basic APPRIS P1	
Dnajb13-201	ENSMUST0000054923.8	847	<u>256aa</u>	Protein coding	e.	<u>A0A171EBK9</u>	CDS 5' incomplete TSL:3	
Dnajb13-203	ENSMUST00000154516.2	905	<u>114aa</u>	Nonsense mediated decay	27	A0A140LHH6	TSL:5	
Dnajb13-202	ENSMUST00000130534.7	1159	No protein	Processed transcript		-	TSL:1	

The strategy is based on the design of *Dnajb13-204* transcript, the transcription is shown below:

< Dnajb13-204 protein coding

Reverse strand -

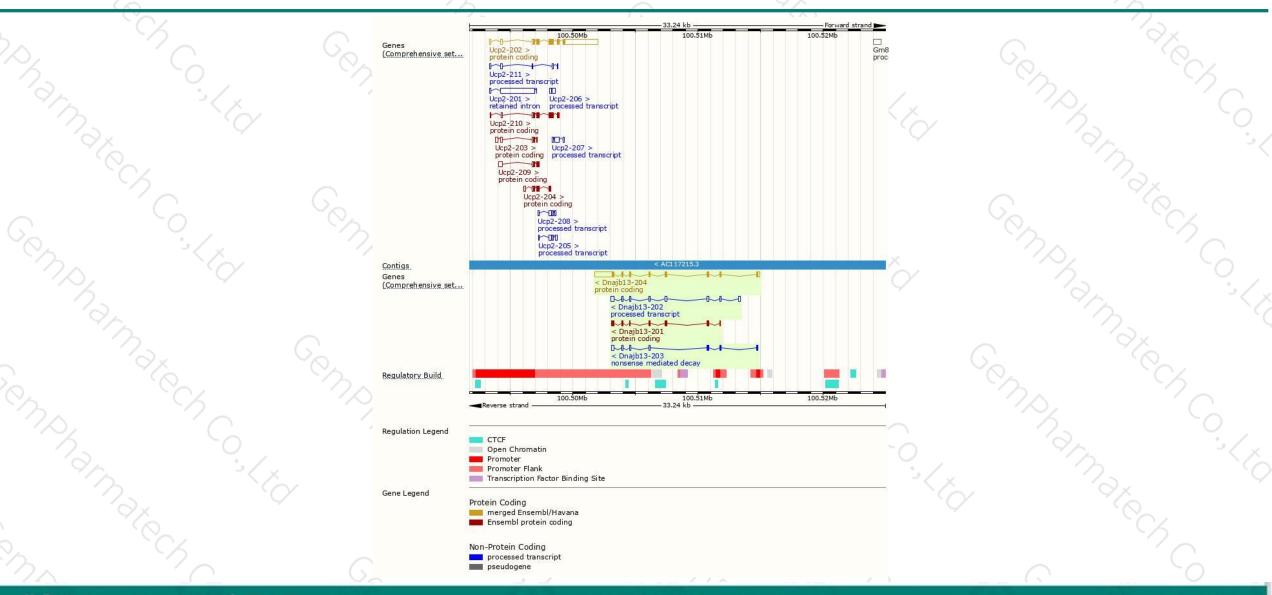
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13.24 kb

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Genomic location distribution



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Protein domain



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	Prints	DnaJ domain	_						
	Pfam	DnaJ domain		c	haperone DnaJ, C	-terminal			°-<
	PROSITE profiles	DnaJ domain	-						
G.	PROSITE patterns	DnaJ d	omain, conserved s	ite					
~?;	PANTHER	PTHR24078:SF519							
•		PTHR24078						-	-34
	Gene3D	Chaperone J-domain supe	erfamily	2.6	0.260.20	-			
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If you have any questions, you are welcome to inquire. Tel: 400-9660890



