

Cd7 Cas9-CKO Strategy

Designer: Reviewer: Design Date:

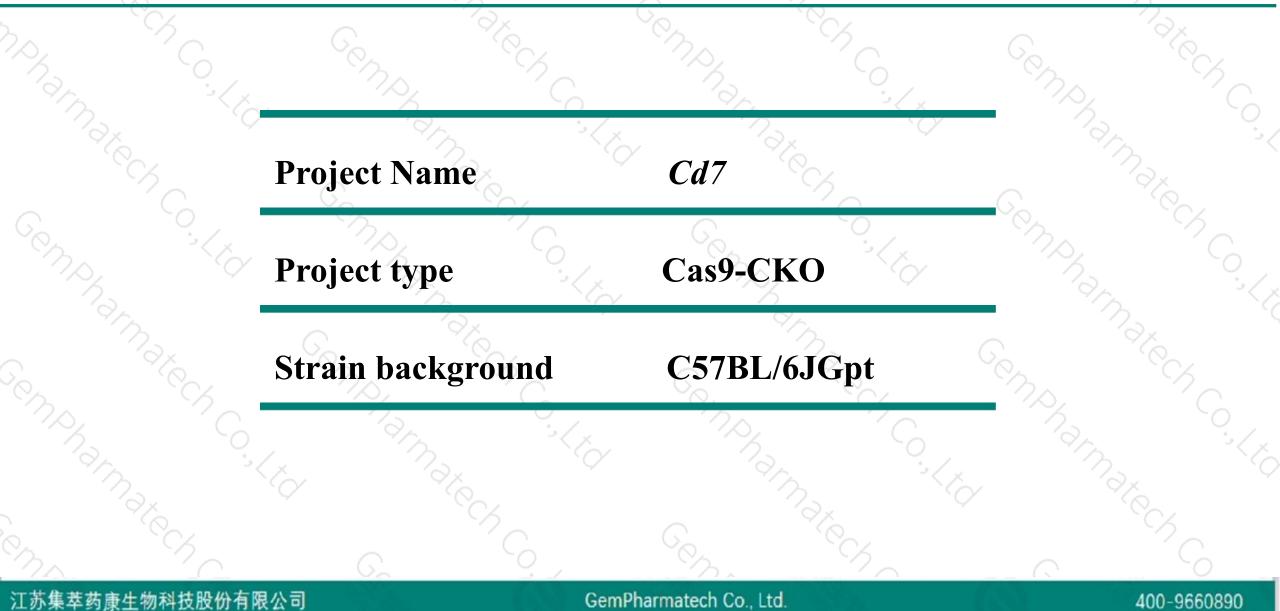
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Project Overview



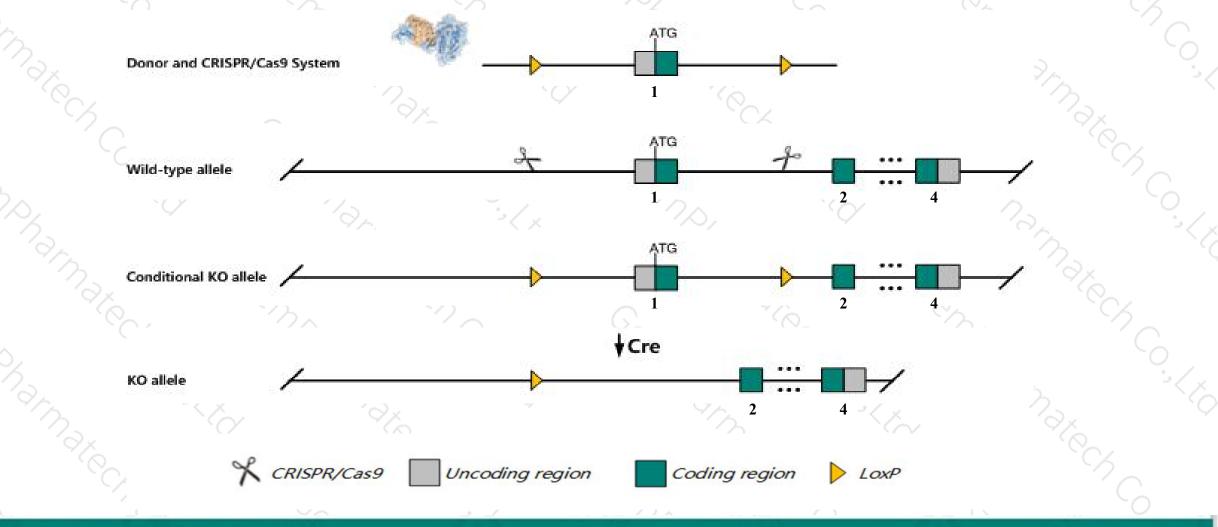


Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the Cd7 gene. The schematic diagram is as follows:



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The Cd7 gene has 2 transcripts. According to the structure of Cd7 gene, exon1 of Cd7-201 (ENSMUST00000026159.5) 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 Cd7 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



According to the existing MGI data, homozygous mutation of this gene results in a 60% increase of CD4+CD8+ thymocytes at 3 months of age.

The KO region contains intron of the *Gm11775* gene. Knockout the region may affect the function of *Gm11775* gene.

The Cd7 gene is located on the Chr11. 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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Cd7 CD7 antigen [Mus musculus (house mouse)]

Gene ID: 12516, updated on 12-Aug-2019

Summary

Official Symbol	Cd7 provided by MGI								
Official Full Name	CD7 antigen provided by MGI								
Primary source	MGI:MGI:88344								
See related	Ensembl:ENSMUSG0000025163								
Gene type	protein coding								
RefSeq status	VALIDATED								
Organism	Mus musculus								
Lineage	e Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae;								
	Murinae; Mus; Mus								
Expression	Biased expression in spleen adult (RPKM 29.9), duodenum adult (RPKM 18.7) and 7 other tissues See more								
Orthologs	human all								
7. Jr.	$\frac{1}{2}$								



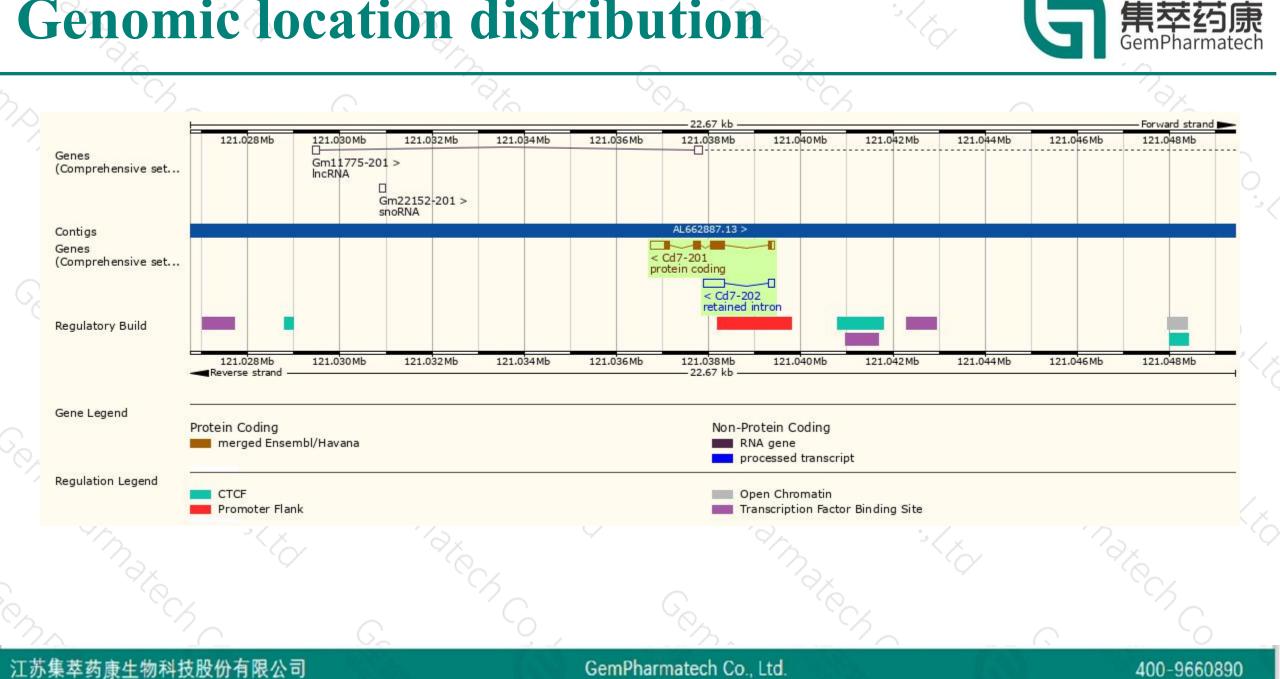
The gene has 2 transcripts, all transcripts are shown below:

Name 🖕	Transcript ID	bp 🖕	Protein 🖕	Biotype 🍦	CCDS 🍦	UniProt 🖕	Flags 🍦	
Cd7-201	ENSMUST0000026159.5	982	<u>210aa</u>	Protein coding	<u>CCDS25764</u> &	P50283 & Q3U4A8 &	TSL:1 GENCODE basic APPRIS P1	
Cd7-202	ENSMUST00000163465.1	592	No protein	Retained intron	-		TSL:2	

The strategy is based on the design of *Cd7-201* transcript, the transcription is shown below:

< Cd7-201 protein coding							
Reverse stra			2.67	kb			
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Genomic location distribution



Protein domain

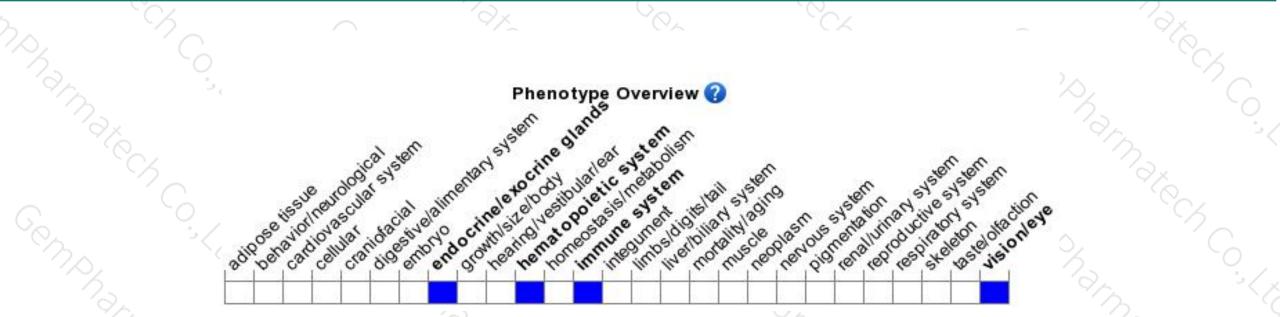
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	Destruction									
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	SMART		Immunoglobulin si	ubtype						
			Immunoglob	ulin V-set domain	(
	<u>Pfam</u>	I	nmunoglobulin V-s	set domain		-				
7	PROSITE profiles	Immune	globulin-like domi	ain						
	PANTHER	T-cell antigen CD7	-							-34
	Gene3D	Contraction of the second second	unoglobulin-like fo	d			-			
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	Variant Legend	missense vi splice regio	n variant						<u></u>	
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集	岑药康生物科技股份有限	公司	16	GemPharr	natech Co., Lt	d.		1	400-9660	0890

Mouse phenotype description(MGI)





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

According to the existing MGI data, Homozygous mutation of this gene results in a 60% increase of CD4+CD8+ thymocytes a 3 months of age.



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



