

# *Madcam1* Cas9-CKO Strategy

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# Overview

## Target Gene Name

- *Madcam1*

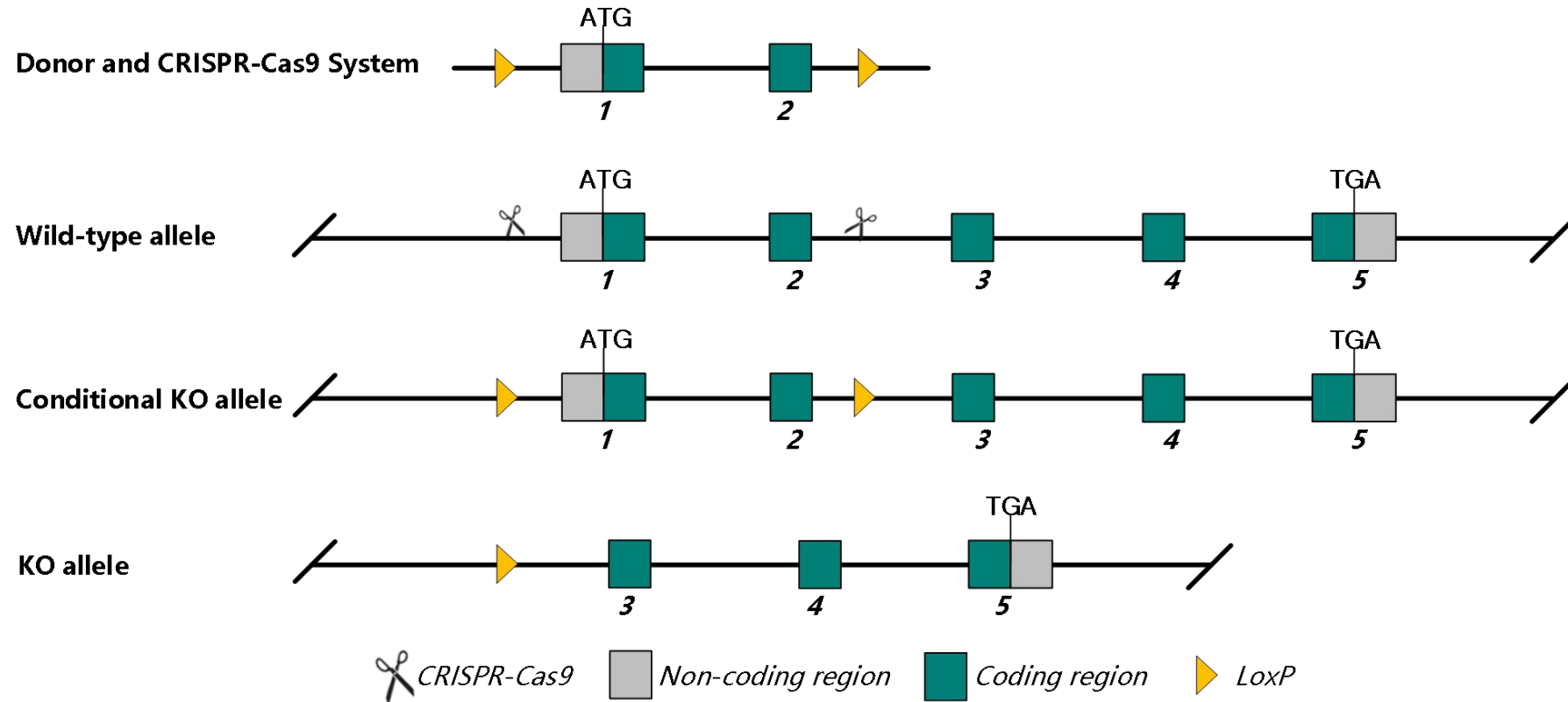
## Project Type

- Cas9-CKO

## Genetic Background

- C57BL/6JGpt

# Strain Strategy



Schematic representation of CRISPR-Cas9 engineering used to edit the *Madcam1* gene.

# Technical Information

- The *Madcam1* gene has 2 transcripts. According to the structure of *Madcam1* gene, exon 1-2 of *Madcam1*-201 (ENSMUST00000020554.8) is recommended as the knockout region. The region contains the start codon ATG. Knocking out the region will result in disruption of gene function.
- In this project we use CRISPR-Cas9 technology to modify *Madcam1* 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 on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.
- 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.

# Gene Information

**Madcam1** mucosal vascular addressin cell adhesion molecule 1 [ *Mus musculus* (house mouse) ]

[Download Datasets](#)

Gene ID: 17123, updated on 23-Nov-2023

## Summary

<b>Official Symbol</b>	Madcam1 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	mucosal vascular addressin cell adhesion molecule 1 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:103579</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000020310</a> <a href="#">AllianceGenome:MGI:103579</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	MAdCAM-1
<b>Summary</b>	Predicted to enable integrin binding activity involved in cell-matrix adhesion. Acts upstream of or within keratinocyte differentiation and leukocyte migration. Predicted to be located in membrane. Predicted to be integral component of membrane. Is expressed in eyelid; hair follicle; and hemolymphoid system. Orthologous to human MADCAM1 (mucosal vascular addressin cell adhesion molecule 1). [provided by Alliance of Genome Resources, Apr 2022]
<b>Expression</b>	Biased expression in spleen adult (RPKM 8.6), large intestine adult (RPKM 4.9) and 7 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>
<b>NEW</b>	Try the new <a href="#">Gene table</a> Try the new <a href="#">Transcript table</a>

## Genomic context

**Location:** 10 C1; 10 39.72 cM

See Madcam1 in [Genome Data Viewer](#)

**Exon count:** 5

<https://www.ncbi.nlm.nih.gov/gene/17123>

# Transcript Information

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

Show/hide columns (1 hidden)							Filter	
Transcript ID	Name	bp	Protein	Biotype	CCDS	UniProt Match	Flags	
<a href="#">ENSMUST00000020554.8</a>	Madcam1-201	1436	<a href="#">405aa</a>	Protein coding	<a href="#">CCDS23981</a>	<a href="#">G5E838</a>	Ensembl Canonical	GENCODE basic APPRIS P4 TSL:1
<a href="#">ENSMUST00000217748.2</a>	Madcam1-202	944	<a href="#">261aa</a>	Protein coding		<a href="#">Q61826-2</a>	GENCODE basic	APPRIS ALT2 TSL:1

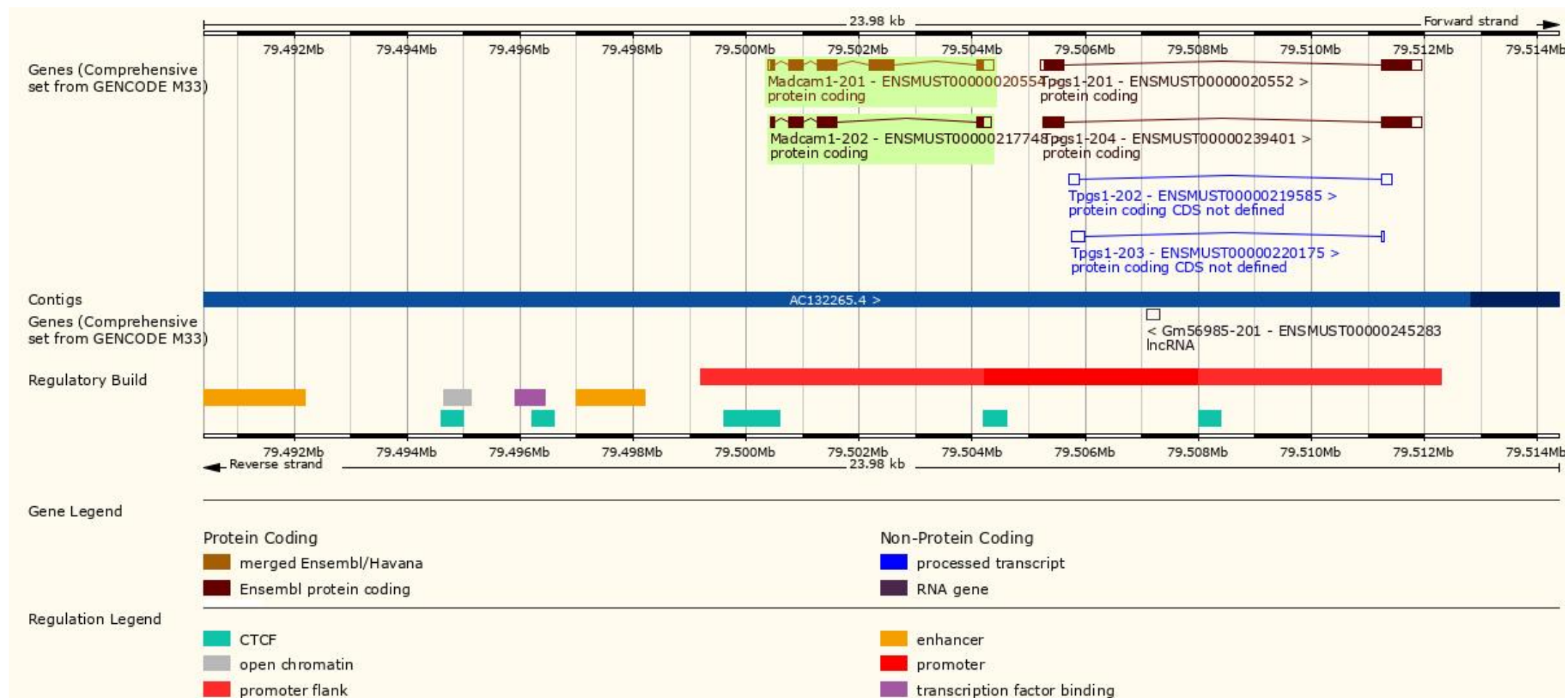
The strategy is based on the design of *Madcam1*-201 transcript, the transcription is shown below:



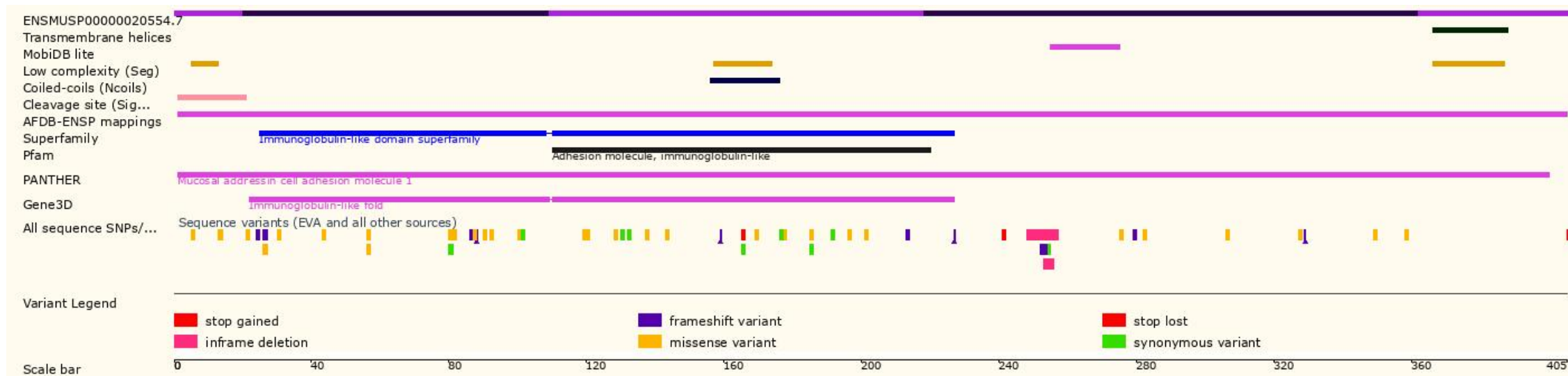
Source: <http://asia.ensembl.org/>



# Genomic Information

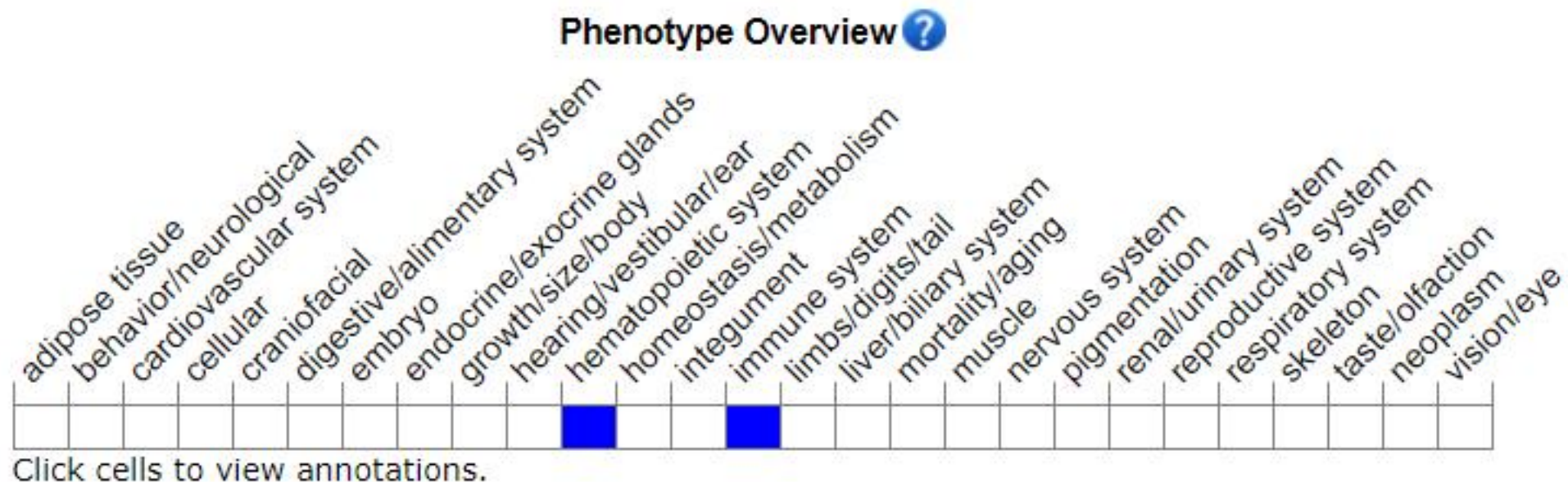


# Protein Information





# Mouse Phenotype Information (MGI)

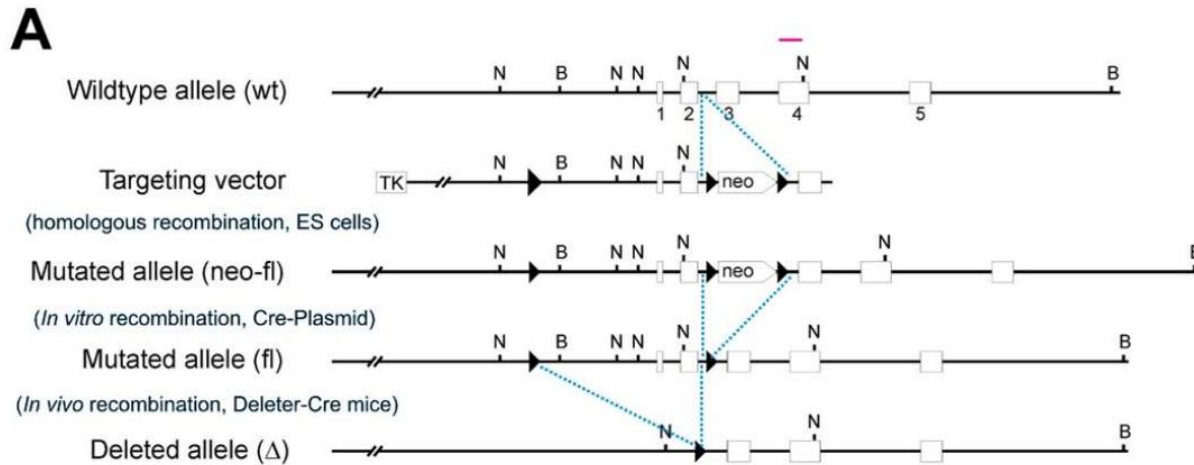


Mice homozygous for a knock-out allele exhibit small Peyer's patches and decreased homing of IgA-secreting plasma cells in the lamina propria.

# Important Information

- The knockout region is about 4 kb away from the 5' of the *Tpgs1* gene, which may affect the regulation of this gene.
- The knockout region contains start codon, translation may recognize new start codon and form new unknown protein.
- The intron 2-3 of *Madcam1* is 252 bp, the loxp insertion may affect the regulation of this gene.
- *Madcam1* is located on Chr 10. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is 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 the existing technology level.

# Reference



## *Deletion of MAdCAM-1 in Mice*

MAdCAM-1-deficient mice (*MAdCAM*<sup>Δ</sup>) were generated by removal of the promoter region and exons 1 and 2 of the *MAdCAM-1* gene (Figure 1A). Complete deletion of the *MAdCAM-1* gene was confirmed by Southern blot analysis (Figure 1B), Northern blot/reverse-transcription polymerase chain reaction (RT-PCR) analysis (Figure 1C, D), and immunofluorescence staining of spleen sections using an anti-MAdCAM-1 antibody (Figure 1E). MAdCAM-1-deficient mice were viable, fertile, and did

[1] Schippers A, Leuker C, Pabst O, Kochut A, Prochnow B, Gruber AD, Leung E, Krissansen GW, Wagner N, Müller W. Mucosal addressin cell-adhesion molecule-1 controls plasma-cell migration and function in the small intestine of mice. *Gastroenterology*. 2009 Sep;137(3):924-33. doi: 10.1053/j.gastro.2009.05.039. Epub 2009 May 18. PMID: 19450594.